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SUMMARIES

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ALIGNMENTS

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RESULT 1
AAB81987
Hanai N,
                                                                                                                                  30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                                                                                                     29-SEP-2000; 2000WO-JP006774
                                                                                                                                                                   05-APR-2001.
                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                         cancer.
                                                                                                                                                                                                              Ganglioside; GD3; complementarity determining region; CDR; antibody;
                                                                                                                                                                                                                           Ganglioside
                                                                                                                                                                                                                                       03-JUL-2001
                                                                                                                                                                               WO200123432-A1.
                                                                                                                                                                                                                                                    AAB81987;
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                                                                                                                       (KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                          Shitara K,
                                                                                                                                                                                                                           GD3
                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                          specific antibody related protein SEQ ID NO:
                                                                                                          Nakamura
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                                                                                                          Niwa
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WPI; 2001-266143/27.

New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent

activity in cancer.

The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention Claim 41; Page 168-172; 183pp; Japanese.

Sequence 582 Α,

Query Match 100.0%; Score 3071; DB 4; Best Local Similarity 100.0%; Pred. No. 2.1e-153; Matches 582; Conservative 0; Mismatches 0; Length 582; Indels 0, Gaps 0

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06-APR-2000; 2000JP-00105088
                                                                                                             29-SEP-2000; 2000WO-JP006774
                                                                                                                                                   WO200123432-A1
                                                                                                                                                                       Synthetic
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                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                     IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT
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type complementation-determining
                                      Shitara K,
                                                                                                                                                                                                   GD3; complementarity
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                                                            HAKKO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
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                                                                                                                                                     TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                                                                                                                                           SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                                                                                                                           LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                                                                                                                                           LYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
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                                                                                                                                                                                                                                                                                                                                                                                                               SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSSA
IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 582
                                                                                                                                                                                         TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
                                                                                                                                                                                                                                             SVFLFPFKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                                                                                                                                                                                                                                  SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSSA
                                                                                      QGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGINNY
                                                                                                                                       TKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWO
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Pred. No. 4.9e-151;
5; Mismatches 5;
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RESULT 3
AABB3156
ID AABB
XX AABB
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            Unidentified
                                                                Ganglioside
                                      Ganglioside;
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                                                                                                                    AAB83156;
                                                                                                                                             AAB83156 standard;
                                                                  GM2
                                       GM2;
                                                                                          (first
                                                                antibody-related
                                      antibody; cytostatic; cytotoxic;
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IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT

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RESULT 4
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Best Local Sim:
Matches 526;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 583 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibodies against ganglioside GM2 combined with drugs, radioisotopes or proteins for treatment and diagnosis of cancer.
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                                                                                                              NNYKNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISN
                                                                                                                                                             RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGI
                                                                                                                                                                                                                DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                                                                                                                                                SSGLYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
                                                                                                                                                                                                                                                                                                                                                                                                               SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQ
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                                                            INVIVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 582
                                                                                                                                               RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGI
                                                                                                                                                                                               DELTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVDKS
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                                                                                               NNYKNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLI
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Pred. No. 3.9e-138;
2; Mismatches 32;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an antibody, which can react specif with ganglioside GD2, and is transplanted with a human type complementation-determining domain (CDR), or its fragments. The ant and its derivatives are useful in diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein us the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g. tumors, activity in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human type complementation-determining domain transplanted antibody and derivatives against ganglioside GD2, useful in diagnosis and therapy of e.g. tumors, has low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ganglioside; GD2; mouse; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                         TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
                                                                                            SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                          SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                             LYSLSSVVTVPSSSSLGTQTYICNVNHKPSNTKVDKKVBPKSCDKTHTCPPCPAPELLGGP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shitara K,
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                                                                                                                                                                                                                                                                                                                                                             89.9%; ilarity 90.0%; Conservative 2
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Pred. No. 4.3e-137;
2; Mismatches 35;
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Best Local Similarity
Matches 516; Conserv
                                                                                                                                                                                          The present invention relates to novel recombinant anti-EpcAM (human epithelial cell adhesion molecule) antibodies comprising an amino acid sequence defining an immunoglobulin light or heavy chain framework region. Sequences of the present invention are useful for the diagnosis, prognosis and treatment of cancer. They are also used in gene therapy. The present sequence is KS antibody heavy chain-interleukin 2 (IL-2) fusion protein. This sequence is used to illustrate the method of the
                                                                                                                                                       Sequence 579
                                                                                                                                                                                                                                                                                                                    New recombinant anti-EpCAM antibody having an amino acid sequence defining an immunoglobulin light or heavy chain framework region, for the diagnosis, prognosis and treatment of cancer.
                                                                                                                                                                                tusion protein. invention
                                                                                                                                                                                                                                                                                              Disclosure; Page 80-82; 82pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunoglobulin; diagnosis; epithelial cell adhesion molecule;
cancer; gene therapy; interleukin-2; IL2; fusion protein.
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                   SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR-VKLGTYYFDSWGQGTLLTVSS
                                                  IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT
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IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 581
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88.5%; Pred. No. 1.4e-136;
tive 32; Mismatches 30;
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                                                 New fusion protein comprising a non-IL-2 moiety moiety, useful for preparing a composition for tinfections or immune disorders.
                                                                                                                               Gillies
                                                                                                                                                                               04-DEC-2001;
12-APR-2002;
                                                                                                                                                                                                                      04-DEC-2002; 2002WO-US038780
                                                                                                                                                                                                                                                                                                             Homo sapiens.
Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                         dI-KS-ala-IL2
                                                                                                                                                                                                                                                                                                                                                                                                                   22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAO30910 standard;
                                                                                                    WPI; 2003-513757/48.
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                                                                                                                                                                                                                                                                         WO2003048334-A2
                                                                                                                                                                                                                                                                                                                                                                Cytokine;
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                                                                                                                                                                                                                                                                                                                                                  therapy;
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                                                                                                                                                                                                                                                                                                                                                                interleukin-2;
                                                                                                                                                        LEXIGEN
                                                                                                                                                                               2001US-0337113P.
2002US-0371966P.
                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                  immunoglobulin; Ig;
                                                                                                                                                                                                                                                                                                                                                                                         (D20T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                        RES
                                                                                                                                                                                                                                                                                                                                                                                        variant
                                                                                                                                                          CENT
                                                                                                                                                                                                                                                                                                                                                                IL-2;
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                                                                                                                                                          CORP
                                                                                                                                                                                                                                                                                                                                                                                        protein.
                                                                                                                                                                                                                                                                                                                                                 cancer; viral infection; immune disorder;
Ig; fusion protein; human.
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                                                              reating cancer, viral
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Example 10; Page 60-63;

71pp; English.

invention relates to cytokine fusion proteins with increased

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RESULT 7
ADD42961
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Matches
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            WO2004055056-A1
                                                                immunoglobulin;
                                                                                       Humanised
                                                                                                                                                       ADP42961
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                                                                                     immunoglobulin heavy chain-IL-2 fusion
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                                                       <u>Burface</u>
                                                                                                          (first
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                                                     variable region; antibody; GD2;
rface glycosphingolipid; IL-2.
                                                                                                                                                      protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               89.6%;
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Pred. No. 1.4e-136;
2; Mismatches 30;
                                                                cytostatic;
                                                                                    protein SEQ ID NO:6.
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Query Match
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Matches 518
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 6; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New modified m14.18 antibodies with reduced immunogenicity specifically bind the human cell surface glycosphingolipid for treating cancer.
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LYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKRVBPKSCDKTHTCPPCPAPELLGGP
                                                           KNPKLTRMLTFKFYMPKKATELKHLQCLEBELKPLEBVLNLAQSKNFHLRPRDLISNINV
                                                                                                                                       QGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGINNY
                                                                                                                                                                                              TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                                                                                                     TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
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                                                                                                                                                                                                                                                                                               TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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                                      KNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISNINV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.6%; ilarity 89.0%; Conservative 2
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4; Mismatches 33;
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                                                                                                                APTSSSTKKTQLQLEHLLLDLQMILNGINNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8;
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582
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RESULT 8
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Best Local S
Matches 500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New fusion protein comprising a non-IL-2 moiety fused to a mutant IL-2 moiety, useful for preparing a composition for treating cancer, viral infections or immune disorders.
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12-APR-2002;
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                                                                                                                                                                                                                                                                                                                Sequence 580
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                                                                                                                                                                                                                                                                     Local Similarity
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                             178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10; Page 68-71; 71pp; English.
                                                                                                   ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                                                                                                     EVQIVESGGDFVQPGGSIRVSCAASGFAFSH-YAMSWVRQAPGKGIEWVAYISSGGSGTY
 PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                            GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                   ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                           YNPSLKSRVTISVDTSKNOFSLKLSSVTAADTAVYYCARGKWSK--FDYWGQGTLVTVSS
                                                                                                                                                                      YSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                                                                                                                                                                                QVQLQESGPGLVKPSETLSLTCAVSGYSISSGYYWGWIRQPPGKGLEWIGSIYHSGS-TY
                           GLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVDKRVEPKSCDKTHTCPPCPAPEFLGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (gamma4h)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interleukin-2; IL-2; cancer; viral infection; immune disorder;
                                                                                                                                                                                                                                                        85.3%; ilarity 85.8%; Conservative 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LEXIGEN
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2002US-0371966P.
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                                                                                                                                                                                                                                                        29;
                                                                                                                                                                                                                                                      Score 2621; DI
Pred. No. 9.2e
29; Mismatches
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                                                                                                                                                                                                                                                                     DB 6;
.2e-130;
                                                                                                                                                                                                                                                                                  Length 580;
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RESULT 9
AAOJ0913
AD AAOJ
XX AAOJ
AC AAOJ
XX AOJ
AC AAOJ
XX Qene
XX Cyto
XX Gyto
XX Gyto
XX Homo
OS Unid
XX 12-J
XX 12
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                                                The invention relates to cytokine fusion proteins with increased therapeutic index and methods for increasing the therapeutic index of such fusion proteins. The fusion protein comprises a non-interleukin-2 (IL-2) moiety fused to a mutant IL-2 moiety. It is useful for preparing a composition for treating cancer, viral infections or immune disorders. The fusion protein is also used in gene therapy. The present sequence is dI-NHS76 (gammazh) (FN>AQ)-ala-IL2 (D2OT) variant protein comprising dI-XS heavy chain fused to human IL-2 (D2OT) variant protein. This sequence is used to illustrate the method of the invention
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12-APR-2002;
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  Sequence 580
                                                                                                                                                                                                                                                                                                                                                                                                                New fusion protein comprising a non-IL-2 moiety fused to a mutant IL-2 moiety, useful for preparing a composition for treating cancer, viral
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                                                                                                                                                                                                                                                                                                                                                                                            infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-513757/48
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Unidentified.
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apy; immunoglobulin; ]
                                                                                                                                                                                                                                                                                                                                                                                               or immune
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                                                                                                                                                                                                                                                                                                                                                                                               disorders.
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                                                                                                                                                                                                                                                                                                                                     English.
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fibrosis), allergic rhinitis, respiratory distress syndrome, asthma, bronchitis, tendonitis, bursitis, fever, migraine headaches, gastro-
intestinal conditions (e.g. inflammatory bowel disease, Crohn's disease, gastritis, irritable bowel syndrome, colitis and colorectal cancer), vascular diseases (e.g. atherosclerosis), thyroiditis, aplastic anaemia, periarteritis nodosa, Hodgkin's disease, rheumatic fever, osteoarthritis, autoimmune diseases (e.g. type I diabetes, myasthenia gravis, rheumatoid arthritis, systemic lupus erythematosus and multiple sclerosis), renal failure, sarcoidosis, nephrotic syndrome, Behcet's syndrome, gingivitis, polymyositis, hypersensitivity (e.g. delayed type hypersensitivity or immediate hypersensitivity), graft and transplant rejections, graft versus host disease, conjunctivitis, swelling occurring after injury, myocardial ischaemia or endotoxin shock syndrome. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to novel antibodies that specifically bind to very late activation (VLA-1; betal containing integrins) antigens and methods of using these antibodies to treat immunological disorders. The anti-VLA-1 antibodies are useful for preventing or treating VLA-1-mediated immunological or inflammatory disorders such as skin related conditions (e.g. psoriasis, eczema, burns, dermatitis and abnormal proliferation of hair follicle cells), fibrosis (e.g. kidney or lung the proliferation of hair follicle cells), fibrosis (e.g. kidney or lung the proliferation of hair follicle cells), fibrosis (e.g. kidney or lung the proliferation of hair follicle cells).
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                                                                       SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                 LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                         STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPBPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                                                                                LDSVKGRFTISRDNSKNTLYLQMNSLRABDTAVYYCTRGFGDGGYFDVMGQGTLVTVSSA
                                                                                                                                                                                                                                                             SDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSSA
                                                                                                                                                                                                                                                                                       EVQLVESGGGLVQPGGSLRLSCAASGFTFSRYIMSWVRQAPGKGLEWVATISGGGH-TYY
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                                                        SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                        STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
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95.1%;
                                                                                                                                                                                                                                                                                                                                                  Score 2246.5; DB 6;
Pred. No. 3.3e-110;
4; Mismatches 17;
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WO200283854-A2

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RESULT 11
AAE33523
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                 to very late activation (VLA-1; betal containing integring) antigens and methods of using these antibodies to treat immunological disorders. The anti-VLA-1 antibodies are useful for preventing or treating VLA-1 mediated immunological or inflammatory disorders such as skin related conditions (e.g. psoriasis, eczema, burns, dermatitis and abnormal proliferation of hair follicle cells), fibrosis (e.g. kidney or lung fibrosis), allergic rhinitis, respiratory distress syndrome, asthma, bronchitis, tendonitis, bursitis, fever, migraine headaches, gastrointestinal conditions (e.g. inflammatory bowel disease, Crohn's disease, gastritis, irritable bowel syndrome, colitis and colorectal cancer), vascular diseases (e.g. atherosclerosis), thyroiditis, aplastic anaemia, periarteritis nodosa, Hodgkin's disease, myasthenia gravis, rheumatoid autoimmune diseases (e.g. type I diabetes, myasthenia gravis, rheumatoid faither servicionis periarters systemic lupus erythematosus and multiple earliers; rheumatoid in the colorest of the colorest 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New anti-VLA-1 antibodies are useful for preventing or treating VLA-1-mediated immunological or inflammatory disorders, e.g. psoriasis, eczemediated immunological or inflammatory disorders, e.g. psoriasis, eczemediated immunological or inflammatory disorders, e.g. psoriasis, dermatitis, and abnormal proliferation of hair follicle cells o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunological disorder; inflammatory disorder; skin related condition; psoriasis; eczema; burn; dermatitis; respiratory distress syndrome; fibrosis; allergic rhinitis; asthma; bronchitis; tendonitis; bursitis; fever; migraine headache; inflammatory bowel disease; Crohn's disease; irritable bowel syndrome; colitis; colorectal cancer; vascular disease; atherosclerosis; thyroiditis; aplastic anaemia; periarteritis nodosa; gastritis; Hodgkin's disease; rheumatic fever; autoimmune disease; osteoarthritis; type I diabetes; myasthenia gravis; rheumatoid arthritis; systemic lupus erythematosus; multiple sclerosis; nephrotic syndrome; renal failure; sarcoidosis; Behcet's syndrome; gingivitis; polymyositis; hypersensitivity; graft rejection; transplant rejection; conjunctivitis; graft rejection; transplant rejection; conjunctivitis;
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06-JUL-2001; 2001US-0303689P
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                                                                                                                                                                                                                                                                                                                                                                                                present invention relates to novel antibodies that specifically bind very late activation (VLA-1; betal containing integrins) antigens and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 91-92; 248pp; English
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     Behcet's
gingivitis,
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Best Local Similarity
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              QGNVFSCSVMHEALHNHYTQKSLSLSPG 448
                                                         TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                     TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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QGNVFSCSVMHEALHNHYTQKSLSLSPG
                                                                                         TKNOVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWO
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                                                                                                                     TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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94.9%;
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Pred. No. 6.8e
4; Mismatches
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3.8e-110;
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RESULT 12
AAO18400
13-OCT-2000;
13-MAR-2001;
21-JUN-2001;
                                                                                                     WO200230986-A2
                                                                                                                        Synthetic.
                                                                                                                                                    Mouse; humanised antibody; lymphotoxin beta receptor; antibody; neoplasia; LT-beta-R; light chain; heavy chain; variable region
                                                                                                                                                                                 Mature humanised murine CBE11 heavy chain variable
                                                                                                                                                                                                    11-OCT-2002
                                                                                                                                                                                                                       AAO18400;
                                                                                                                                                                                                                                        AAO18400 standard; protein; 449 AA
                                                               12-OCT-2001; 2001WO-US032140.
                                                                                   L8-APR-2002.
        (BIOJ )
        BIOGEN INC
                          2000US-0240285P.
2001US-0275289P.
2001US-0299987P.
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RESULT 1
AAE33524
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Matches 422
Human; very late activation antigen; VLA-1; betal containing integrin; immunological disorder; inflammatory disorder; skin related condition; psoriasis; eczema; burn; dermatitis; respiratory distress syndrome; fibrosis; allergic rhinitis; asthma; bronchitis; tendonitis; bursitis; fever; migraine headache; inflammatory bowel disease; Crohn's disease; irritable bowel syndrome; colitis; colorectal cancer; vascular disease; atherosclerosis; thyroiditis; aplastic anaemia; periatteritis nodosa; gastritis; Hodgkin's disease; rheumatic fever; autoimmune disease;
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present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New humanized anti-lymphotoxin-beta receptor antibody, useful for treating or reducing the advancement, severity or effects of neoplasia, particularly solid tumors (i.e. carcinomas) including colorectal cancer
                                                                                                                                                                             02-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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(LT-beta-R) antil
pinding antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nction relates to humanised anti-lymphotoxin beta receptor bodies. These are derived from the murine LT-beta-R CBE11 and can be used to treat neoplasia in humans. The is a humanised murine CBE11 heavy chain variable region
                                                                                                                                           chain mutant protein, hsAQC2.
                                                                                                                                                                                                                                        protein;
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systemic lupus erythematosus; multiple sclerosis; nephrotic syndrome; renal failute; sarcoidosis; Behcet s syndrome; gingivitis; polymyositis; hypersensitivity; graft rejection; transplant rejection; conjunctivitis; graft versus host disease; myocardial ischaemia; mutant; mutein.
                                                                                                                                                                                                                                                                            13-APR-2001; 2001US-0283794P 06-JUL-2001; 2001US-0303689P
                                                                                                                                                                                                                                                                                                   12-APR-2002;
                                                                                                                                                                                                                                                                                                                                WO200283854-A2
                                                                                                                                                                                                                                                                                                                                                                                          osteoarthritis; type I diabetes; myasthenia gravis; rheumatoid arthritis,
                                                                                                                                                                                                                                                              (BIOJ ) BIOGEN INC
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mediated anti-VLA-1 antibodies are useful for preventing or treating VLA-1-iated immunological or inflammatory disorders, e.g. psoriasis, eczerns, dermatitis, and abnormal proliferation of hair follicle cells o 23; Page 92; English

Example

WPI; 2003-093009/08

CC to very late activation (VIA-1; betal containing integrins) antigens and CC methods of using these antibodies to treat immunological disorders. The CC anti-VIA-1 antibodies are useful for preventing or treating VIA-1-CC mediated immunological or inflammatory disorders such as skin related CC conditions (e.g. psoriasis, eczema, burns, dermatitis and abnormal CC proliferation of hair follicle cells), fibrosis (e.g. kidney or lung CC fibrosis), allergic rhinitis, respiratory distress syndrome, asthma, CC bronchitis, tendonitis, burshtis, fever, migraine headaches, gastro-CC conditions (e.g. inflammatory bowel disease, Crohn's disease, CC qastritis, irritable bowel syndrome, colitis and colorectal cancer), and colorectal cancer), and colorectal cancer, cC continumnte diseases (e.g. atherosclerosis), thyroiditis, aplastic anaemia, CC periarteritis nodosa, Hodgkin's disease, rheumatic fever, osteoarthritis, CC autoimmune diseases (e.g. type I diabetes, myasthenia gravis, rheumatoid carchritis, systemic lupus erythematosus and multiple sclerosis), renal CC failure, sarcoidosis, nephrotic syndrome, Behcet's syndrome, gingivitis, CC polymyositis, hypersensitivity (e.g. delayed type hypersensitivity or immediate hypersensitivity), graft and transplant rejections, graft consuman AQC2 heavy chain mutant protein, hsAQC2 invention relates to novel antibodies that specifically bind

Sequence 447 A,

Similarity

72.8%;

No. 1.1e

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                       the formation of CD4+ memory cells. A human OX40 receptor specific phage antibody (scFv) represents the agonistic binding molecule of the invention which is capable of binding to and stimulating the human OX-40-receptor. The binding molecules of the invention are useful as a medicament for treating neoplastic, viral or bacterial disorder or disease, for stimulating T-cells in vitro and for enhancing the immune response in a human or animal against a tumour, bacterial or viral antigen. They are useful for modulating T-cell response in a human which involves, administration to the human, where the modulation includes
                                                                                                                                                                                                                                                                                This sequence represents the heavy chain of a selected phage antibody against human OX40 receptor. OX40 receptor. (also known as CD134, ACT-4, ACT35) is a member of the TNF receptor family which is expressed on activated CD4+ T cells. Triggering of this receptor enhances the proliferation of CD4+ T-cells during an immune response and influences the formation of CD4+ memory cells. A human OX40 receptor specific phage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody; OX40 receptor; specific phage antibody; scFv; CD134; ACT-4; ACT35; TNF receptor family; CD4+; T cell; tumour necrosis factor; proliferation; immune response; memory cell; tumour; bacterial; viral; antigen; leukaemia; neoplasm; bile duct carcinoma; bladder carcinoma; severe acute respiratory syndrome; SARS; herpes simplex virus; HSV; hepatitis B virus; HSV; HIV; Mycobacterium tuberculosis; Pseudomonas aeruginosa; Vibrio cholerae.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                           Human A2-G8 SCF antibody heavy chain variable and constant region
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                                                                                                                                                                                                                                                                                                    AAO31101 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVK--LGTYYFDSWGQGTLLTVS
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                                                                                                                                                                                                                                                                                                                                                                                WQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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93.1%;
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Human; antibody; stem cell factor; mast cell growth factor; asthma; steel factor; c-kit ligand; gene therapy.

WO2003051311-A2

26-JUN-2003.

Takeuchi T,

Tomkinson A,

Neben

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17-DEC-2001;

2001US-0342174P

FARB) BAYER CORP

16-DEC-2002; 2002WO-US040227

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Best Local S
Matches 421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides human antibodies that bind to stem cell factor (SCF) protein. SCF is also known as mast cell growth factor, steel factor or c-kit ligand. Antibodies of the invention are useful for preparing compositions for treating asthma. They are also used in gene therapy. The present sequence is human SCF antibody heavy chain variable and constant
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 445 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New purified human antibody that binds to stem cell factor protein, useful for preparing a composition for treating asthma.
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                                               424
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             VFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                     LFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYR 303
                                                                                                                                                                                                   LSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVF 243
                                                                                                                                                                                                                                                                                                     VKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSSASTK 123
                                                                                                                                                    LFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYR
                                                                                                                                                                                                                                                      GPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYS
                                                                                                                                                                                                                                                                       GPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYS 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 47; 94pp; English.
                                                                                                   VVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKN
VFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                                                                                                                                                                                                                            72.8%; ilarity 94.4%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          Score 2234.5; DB 6;
Pred. No. 1.4e-109;
7; Mismatches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 445;
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Search completed: December 23, 2004, 18:57:36 Job time : 160.692 secs

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Sequence 2, Appli Sequence 1, Appli Sequence 18, Appli Sequence 18, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 8, Appli Sequence 8, Appli Sequence 10, Appli Sequence 113, Appli Sequence 12, Appli Sequence 12, Appli Sequence 14, Appli Sequence 41, Appli

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             seq length: 0
seq length: 2000000000
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Match
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3071
1 EVQLVESGGDFVQPGGSLRV.....IVEFLNRWITFCQSIISTLT
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1: /cgn2_6/ptodata/1/laa/5A_COMB.pep:*
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3: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*
4: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/laa/PCTUS_COMB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                 Length
    GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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US-09-026-985-71
US-09-121-9528-71
US-09-234-340A-71
US-09-121-9528-19
US-08-466-1538-8
US-08-802-096-8
US-08-802-097-8
US-08-807-3528-14
US-08-807-3528-14
US-08-807-3528-14
US-08-807-3528-15
US-08-96-109-207C-14
US-09-109-207C-14
US-09-296-005-16
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US-09-296-019-10-14
US-09-296-019-10-14
US-09-296-019-10-14
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US-09-296-019-16
US-09-296-019-16
US-09-296-019-16
US-09-109-028-14
US-09-113-996-16
US-10-113-996-16
US-10-113-996-16
US-09-282-505-2
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                   -282-505-2
-054-255-2
Sequence 71, Appl
Sequence 8, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 14, Appli
Sequence 16, Appli
Sequence 16, Appl
Sequence 11, Appl
Sequence 12, Appl
Sequence 13, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 18, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                   Description
                                                          ; SEQUENCE CHARACTERISTICS:
; LENGTH: 452 amino acids
; TYPE: Amino Acid
; TOPOLOGY: Linear
US-09-027-449-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1
US-09-027-449-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 71, Application US/09027449
Patent No. 6025158
GENERAL INFORMATION:
   Query Match 71.5%;
Best Local Similarity 90.9%;
Matches 411; Conservative 1.
                                                                                                                                                                     COMPUTER: IBM PC COMPALIBLE
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/027,449
FILING DATE: 20-Feb-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/074,330
FILING DATE: 22-Jan-1998
PRIOR APPLICATION NUMBER: 60/038,664
FILING DATE: 21-Feb-1997
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: p1085R3-2
TELEPHONE: 60/025,5530
                                                                                                                                 TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibo
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1 DNA Way
CITY: South San Fr
STATE: California
                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Francisco
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US-09-20-171-18
US-09-20-171-18
US-09-716-028-18
US-09-483-588-2
US-10-113-996-18
US-09-679-97-2
US-09-680-148-2
US-09-304-465A-2
US-09-304-465A-2
US-09-383-916-8
US-09-383-916-8
US-09-383-916-13
US-08-485-516-13
US-08-485-7378-939-10
                                                                                                                                                                                             P1085R3-2
   Score 2194.5; DB 3;
Pred. No. 2.3e-161;
4; Mismatches 24;
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                              Length
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Antibodies
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Result No.

Score

452; <u>ب</u>

Minimum DB Maximum DB

Scoring table: Sequence: Perfect score: Run on:

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STREET: 1 DAN WAY
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy dis
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winpatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/026,985
FILING DATE: 20-Feb-1998
CLASSIFICATION:
NAME: LOVE, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: 91085R3-1
TELEPAN: 650/25-5530
TELEPAN: 650/25-9881
INFORMATION FOR SEQ ID NO: 71:
CEPTIENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2
US-09-026-985-71
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Patent No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Gonzalez, Tania R.
APPLICANT: Gonzalez, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates and
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibodies
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
SEQUENCE CHARACTERISTICS:
LENGTH: 452 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          421
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APPLICANT: Fresta, Leonard G.

APPLICANT: Shahrakh, Zahra

APPLICANT: Zapara, Gerardo A.

TITLE OF INVENTION: METHODS OF TREATING INFLAMMATORY DISEASES

TITLE OF INVENTION: MIH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES

NUMBER OF SEQUENCES: 72

CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
TTP: South San Francisco
COUNTRY: South San Francisco
COUNTRY: USA
ZIPP
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US-09-121-952A-71
; Sequence 71, Applic
; Patent No. 6458355
; Patent No. 6458355
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Best Local S
Matches 411
                                         ZIP: 94080

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/121,952A

FILING DATE: 24-Jul-1998

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA
             APPLICATION NUMBER: 60/0 FILING DATE: 22-JAN-1998
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Sequence 71, Application US/09234340A
Patent No. 6468532
PARPLICANT: Genentech, Inc., Hsei, V
APPLICANT: Koumenis, Iphigenia
APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; SEQUENCE CHARACTERISTICS:
; LENGTH: 452 amino acids
; TYPE: Amino Acid
; TOPOLOGY: Linear
US-09-121-952A-71
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FILING DATE: 20.FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: p1085R
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/252-5530
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 71:
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Best Local S
Matches 411
                                                                                            APPLICANT: Prest, Leonard G.
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
APPLICANT: Zapata, Gerardo A.
TITLE OF INVENTION: METHODS OF TREATING INFLAMMATORY DISEASES
TITLE OF INVENTION: WITH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES
NUMBER OF SEQUENCES: 72
                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech
COUNTRY:
            CITY: South San Francisco
STATE: California
                                               STREET:
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                                               1 DNA Way
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Koumenis, Iphigenia
Leong, Steven R.
Presta, Leonard G.
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                                                           Genentech, Inc.
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Best Local S
Matches 411
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 452 amino acid
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/121,9:

FILING DATE: 24-Jul-1998

APPLICATION NUMBER: 60/074330

FILING DATE: 22-JAN-1998

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/075467

FILING DATE: 20-FEB-1998

ATTORNEY/AGENT INFORMATION:

NAME: LOVE, RICHARD B.

REGISTRATION NUMBER: 34,659

REFERENCE/DOCKET NUMBER: P1085R4

TELECOMMUNICATION INFORMATION:

TELEPHONE: 650/225-5530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Wilphatin (Genentech)
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 71.5%; Score 2194.5; DB 4; Similarity 90.9%; Pred. No. 2.3e-161; 11; Conservative 14; Mismatches 24;
                         RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                           EEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                       DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                                YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                                                                                           GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                                                                                                                                                                                                                  GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                                                                                                                                                                                                                                                                                              SSGLYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL 237
                                                                                                                                                                                                                                                                                                                                                                                                       SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQ
                                                                                                                                                                                                                                                                                                                                                                                                                            SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPBPVTVSWNSGALTSGVHTFPAVLQ 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKL----GTYYFDSWGQGTLLTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EVQLVQSGGGLVQPGGSLRLSCAASGYSFSSHYMHWVRQAPGKGLEWVGYIDPSNGETTY
RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                           YNSTYRVVSVLTVLHODWLNGKEYKCKVSNKALPAPIEKTISKAKGOPREPOVYTLPPSR
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RESULT 5
US-08-157-101A-7
; Sequence 7, Application
; Patent No. 5808032

US/08157101A

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APPLICANT: KURLHARA, TATSUYA
APPLICANT: MATSUKURA, SHIGEKAZU
APPLICANT: TSURUOKA, NOBUO
APPLICANT: ARIMA, KENJI
APPLICANT: NISHHHARA, TATSURO
APPLICANT: NISHHHARA, TATSURO
TITLE OF INVENTION: ANTI-HBS ANTIBODY GE
TITLE OF INVENTION: PLASMIDS THEREFOR
TITLE OF INVENTION: PLASMIDS THEREFOR
OWNEER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 6714627 CUCH
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 459 amino acids
TYPE: amino acid
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NAME: TITUS, MARLANA K
REGISTRATION NUMBER: 35843
REFERENCE/DOCKET NUMBER: 9437
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-861-3711
TELEPAX: 202-822-0944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0,
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TOPOLOGY: lin
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 370
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                  LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
                                                                                        STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                                             PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                                                                                                                                                                                               GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                                                                                                                                                                     GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                                                                                                                                                                                                                   ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALASGVHTFPAVLQSS
                                                                    STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                        PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
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                                                                                                                                                                                                                                                                                                           ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS 179
LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
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91.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 2194.5; DB 1;
Pred. No. 2.3e-161;
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US-08-466-151-8
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                                                                                                                                                                       Query Match
Best Local Similarity 91.01
413; Conservative
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Patent No. 6037453
GENERAL INFORMATION:
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb fl.

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: Winpatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION UMMBER: US/08/466,151

PILING DATE:

PILING DATE:
                                                                                                                                                                                                                                                                                                                                              TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/744
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/405
FILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/18:
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/87:
APPLICATION NUMBER: 07/87:
                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 39,044
REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
TELESPHONE: 460/007
                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 06-Jun-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                          LENGTH: 453 amino TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Svoboda, Craig G
REGISTRATION NUMBER: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U
ZIP: 94080
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SSAST--KGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEFVTVSMNSGALTSGVHTFPAV
                                                                        YSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVK--LGTYYFDSWGQGTLLTV
                                                                                                                 EVQLVESGGGLVQPGGSLRLSCAVSGYSITSGYSWNWIRQAPGKGLEWVASITYDGS-TN
                                                                                                                                       EVQLVESGGDFVQPGGSLRVSCAASGFAF-SHYAMSWVRQAPGKGLEWVAYISSGGSGTY 59
                                                   YADSVKGRFTISRDDSKNTFYLOMNSLRAEDTAVYYCARGSHYFGHWHFAVWGQGTLVTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Francisco
                                                                                                                                                                                        70.6%; Score 2169; DB 3; 91.0%; Pred. No. 2.1e-159; rative 16; Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunoglobulin Variants
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                                                                                                                                                                                                                       Length 453;
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APPLICANT: Jardieu, Paula M.
APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE DE INVENTION: Immunoglobulin Variants
FILE REFERENCE: P0718P2C1D1
CURRENT APPLICATION NUMBER: US/08/466,163B
CURRENT FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR APPLICATION NUMBER: US 08/105,899
PRIOR FILING DATE: 1994-01-26
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
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US-08-466-163B-8
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LENGTH: 453
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial sequence FEATURE:
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                                                                                 LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
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                                                           LOSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
                                                                                                                                                                                                            YADSVKGRFTISRDDSKNTFYLOMNSLRAEDTAVYYCARGSHYFGHWHFAVWGQGTLVTV
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Pred. No. 2.1e-159;
6; Mismatches 19;
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APPLICANT: Jardieu. Paula M.
APPLICANT: Presta, Leonard G.
ITITLE OF INVENTION: Method of Preventing th
ITITLE OF INVENTION: Method of Preventing th
ITITLE OF INVENTION: Method of Preventing th
ITITLE OF INVENTION NUMBER: US/09/802,096
CURRENT FILING DATE: 2001-03-08
PRIOR PELICATION NUMBER: US 08/405,617
PRIOR FILING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
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Best Local Similarity
Matches 413; Conserva
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SEQ ID NO 8
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TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
OTHER INFORMATION: humanized maell, version 1 heavy chain
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RESULT 9
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CURRENT FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR FILING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR PILING DATE: 1994-01-26
PRIOR PPLICATION NUMBER: US 07/US92/06860
PRIOR PPLICATION NUMBER: US 07/879,495
PRIOR PILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR PILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR PILING DATE: 1991-08-14
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Best Local S
Matches 413
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APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Treating Allergic Disorders (as
FILE REFERENCE: P0718P2C2US
FILE REFERENCE: P0718P2C2US
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SEQ ID NO 8
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OTHER INFORMATION: humanized maell, version 1 heavy chain
-09-802-077-8
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TYPE: PRT
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TELECOMENCE: 650/22-9881
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 451 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
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US-08-887-352B-14
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Patent No. 5994511
GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula
TITLE OF INVENTION: Improved Anti-1gE Antibodies and
TITLE OF INVENTION: Improving Polypeptides
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 411; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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MEDIUM TYPE: 3.5 inch, 1.44 Mb flc

MEDIUM TYPE: 3.5 inch, 1.44 Mb flc

MEDIUM TYPE: 3.5 inch, 1.44 Mb flc

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION UNMBER: US/08/887,352E

FILING DATE: 03-Jul-1997

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: Syoboda, Craig G.

REGERENCE/DOCKET NUMBER: 39,044

REFERENCE/DOCKET NUMBER: P1123

TELEPHONE: 650/225-1489
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                  DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                           GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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                                                                    YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                      YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                         GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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90.9%; Pred. No. 3e
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US-08-887-352B-16
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Patent No. 5994511
GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leona
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/887,352B
FILING DATE: 03-Unl-1997
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 9123
REFERENCE/DOCKET NUMBER: P1123
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 451 amino acids
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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STREET: South San Fra
CITY: California
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TITLE OF INVENTION:
TITLE OF INVENTION:
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CORRESPONDENCE ADDRESS:
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                YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR 357
                                                                                    GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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                                                                                                                                      SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
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                                                                GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                       16; Mismatches
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US-08-466-151-65
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                                                                                                                                                                                                                                                                                         TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P071
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PILING DATE:
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/4661
APPLICATION NUMBER: 08/4661
APPLICATION NUMBER: 08/4066
PILING DATE: 15-MAR-1995
APPLICATION NUMBER: 08/1856
PILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/8794
PRIOR APPLICATION NUMBER: 07/8794
PRIOR APPLICATION DATA:
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APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
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                                                                                                                                                                                                                                                             TYPE: Amino Acid
                                                                                                                                                                                                                                              TOPOLOGY: Linear
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                                                                                                                                                                  h 70.6%; Score 2167; DB 3 Similarity 90.9%; Pred. No. 3e-159;
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                         EVQLVESGGGLVQPGGSLRLSCAVSGYSITSGYSWNWIRQAPGKGLEWVASITYDGS-TN
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APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polype; PILE REFERENCE: P1123R1 CURRENT APPLICATION NUMBER: US/09/109,207C CURRENT APPLICATION NUMBER: US 60/051,554

PRIOR APPLICATION NUMBER: US 60/051,554

PRIOR FILING DATE: 1997-07-03

NUMBER OF SEQ ID NO 14

LENGTH: 451

TYPE: PRT
ORGANISM: Artificial FEATURE:
NAME/KEY: Artificial
FEATURE:
NAME/KEY: Artificial
COCATION: 1-451

OTHER INFORMATION: Heavy chain sequence derived from MAE11

US-09-109-207C-14
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GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard
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PRIOR FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 16
LENCTH: 451
TYPE: PRT
ORGANISM: Artificial
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US-09-109-207C-16
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APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptide
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FILE REFERENCE: P1123R1
CURRENT APPLICATION NUMBER: US/09/109,207C
CURRENT FILING DATE: 1998-06-30
CURRENT FILING DATE: US 60/051,554
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NAME/KEY: Artificial
LOCATION: 1-451
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Pred. No. 3e-159;
6; Mismatches 2
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GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
FILE CAI INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptides
FILE REFERENCE: P1123C1r
CURRENT ETLING NUMBER: US/09/296,005
CURRENT ETLING DATE: 1999-04-21
EARLIER APPLICATION NUMBER: US 08/887,352
EARLIER APPLICATION NUMBER: US 08/887,352
NUMBER OF SEQ ID NO 14
LENGTH: 451
TYPE: PRT
ORGANISM: Artificial
FEATURE:
NAME/KEY: Artificial
FEATURE:
NAME/KEY: Artificial
COTHER INFORMATION: Heavy chain sequence derived from MAE11
US-09-296-005-14
Search completed: December 23, 2004, 19:07:50 Job time: 40.4376 secs
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Matches 411; Conserv
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                                                                RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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                                                                  451
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Maximum Match 100%
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Perfect score:
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seq length: 2000000000
Published Applications AA:*

1: /cgn2 6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

3: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

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19: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

111110 8 8 7 6 5 4 3 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Regult No.
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89.6 89.6 88.6 88.3 84.3 73.2 72.9 72.9 72.8 72.8	Query Match Length
579 579 580 580 580 447 447 4852 4457 4444	Length
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Sequence 41, Appl Sequence 32, Appl Sequence 6, Appli Sequence 37, Appli Sequence 37, Appli Sequence 4, Appli Sequence 4, Appli Sequence 5, Appli Sequence 5, Appli Sequence 6, Appli	Description

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e 116	US-10-379-392-		71.1	2184.5	42
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71,	-SD		71.5	2194.5	34
42,	US-1	445	•	2195	ω ω
26,	15 US-10-656-769-26		71.6	2198.5	32
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46,	S		•	2202.5	30
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21,				2209.5	28
12,	US-10-487-322-		•	2209.5	27
34			72.0	2212	26
26,	๘		•	2216	25
12,	S	442	•	2216.5	24
12,			•	2216.5	23
12	15 US-10-226-435A-12		72.2	2216.5	22
1,		474	•	2220	21
ω			72.3	2220	20
Sequence 3, Appli	10 US-09-848-832-3		•	2220	19
Sequence 38, Appl			•	N	18
e 39	14 US-10-291-265-395		•	2222.5	17
e 21	15 US-10-463-442-21		•	2226	16
21,	9 US-09-736-371B-21		72.5	2226	15
Sequence 6, Appli	16 US-10-645-215-6	444	•	2231.5	14

ALIGNMENTS

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PRIOR APPLICATION NUMBER: US/10/138,727)

PRIOR APPLICATION NUMBER: US 60/288,564

PRIOR FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 42

SOFTWARE: Patentin version 3.0

SEQ ID NO 41

LENGTH: 579

TYPE: PRT

ORGANISM: Artificial sequence

FEATURE:
OTHER
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                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: heavy chain-IL2 US-10-138-727A-41
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US-10-138-727A-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Susan
                                                                                                                                                                                    Query Match 89.6%; Score 2750.5; Best Local Similarity 88.5%; Pred. No. 8.2e Matches 516; Conservative 32; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 41, Application US/10138727A Publication No. US20030157054A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Susan
TITLE OZION, SUSAN
TITLE OZION, SUSAN
TITLE OZION, SUSAN
TITLE OZION, SUSAN
TITLE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/10/138,727A
CURRENT FILING DATE: 2002-05-03
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                                                                                                                     1 QIQLVQSGAEVKKPGETVKISCKASGYTFTNYGMNWVKQTPGKGLKWMGWINTYTGEPTY 60
                                                                                                                                                                                                          2e-166;
                                                                                                                                                                                      30; Indels
                                                                                                                                                                                                                                DB 14; Length 579;
                                                                                                                                                                                    5; Gaps
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CURRENT APPLICATION NUMBER: US/10/310,719
CURRENT FILING DATE: 2002-12-04
PRIOR APPLICATION NUMBER: 60/337,113
PRIOR FILING DATE: 2001-12-04
PRIOR PILING DATE: 2001-12-04
PRIOR PILING DATE: 2002-04-12
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.1
SEQ ID NO 32
LENGTH: 579
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 2
US-10-310-719-32
Sequence 32, Application US/10310719
Publication No. US20030166163A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Immunocytokines With Modulated
FILE REFERENCE: LEX-020
                                                                                                                                                                                                                       Query Match 89.6
Best Local Similarity 88.5
Matches 516; Conservative
                                                                                                                                                                                                                                                                                           FEATURE:
OTHER INFORMATION: dI-KS-ala IL2 (D20T) heavy chain fused to IL-2 variant-10-310-719-32
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                                                                                                      SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR-VKLGTYYFDSWGQGTLLTVSS
                                                                                                                                                    EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
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                                        ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
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                                                                                                                                                                                                                      89.6%; Score 2750.5; DB 1
88.5%; Pred. No. 8.2e-166;
ative 32; Mismatches 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: LEX-023
; CURRENT APPLICATION NUMBER: US/10/737,208A
; CURRENT FILING DATE: 2003-12-16
; PRIOR APPLICATION NUMBER: US 60/433,945
; PRIOR FILING DATE: 2002-12-17
; PRIOR FILING DATE: 2002-12-17
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 6
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US-10-737-208A-6
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                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                            Matches 518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Gillies, Stephen D.
APPLICANT: Lo, Kin-Ming
TITLE OF INVENTION: IMMUNOCYTOKINE SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         537
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                                                      LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                        SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLIVSSA 120
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SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                      STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
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                                                                                                      STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                        NQKFKGRATLTVDKSTSTAYMHLKSLRSEDTAVYYCVS---GMEY---WGQGTSVTVSSA
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                                   LYSISSVVTVPSSSIGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELIGGP
                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                         88.6%;
                                                                                                                                                                                                                                                                                                         ; Score 2719.5; ; Pred. No. 7.4e 24; Mismatches
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es 33;
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FILE REFERENCE: LEX-020

CURRENT APPLICATION NUMBER: US/10/310,719

CURRENT FILING DATE: 2002-12-04

PRIOR APPLICATION NUMBER: 60/337,113

PRIOR FILING DATE: 2001-12-04

PRIOR APPLICATION NUMBER: 60/371,966

PRIOR FILING DATE: 2002-04-12

NUMBER OF SEQ ID NOS: 37

SOFTWARE: Patentin version 3.1

SEQ ID NO 37

LENGTH: 580

TYPE: PRT

ORGANISM: Artificial Sequence
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US-10-310-719-37
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; OTHER INFORMATION: dI-NHS76(gamma4h)(FN>AQ)-ala-IL2(D20T)
; OTHER INFORMATION: IL-2 variant
US-10-310-719-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 37, Application US/10310719
Publication No. US20030166163A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Immunocytokines With Modulated Selectivity
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 500; Conserv
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                                       PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                                                                                             GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
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                                                                                         GLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVDKRVEPKSCDKTHTCPPCPAPEFLGG
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Pred. No. 1.3e-157;
9; Mismatches 50;
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US-10-310-719-35
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LENGTH: 580
TYPE: PRT
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APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Immunocytokines With Modulated
FILE REFERENCE: LEX-020
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Publication No. US20030166163A1
                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/371,966
PRIOR FILING DATE: 2002-04-12
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/310,719
CURRENT FILING DATE: 2002-12-04
PRIOR APPLICATION NUMBER: 60/337,113
PRIOR FILING DATE: 2001-12-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                    299 NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRD
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QSTFRVVSVLTVVHQDWLNGKEYKCKVSNKGLPAPIEKTISKTKGQPREPQVYTLPPSRE
                                                                        GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQA
                                                                                            GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQY
                                                                                                                                            SGLYSLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTVEPKSCDKTHTCPPCPAPP-VA
                                                                                                                                                             SGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLG
                                                                                                                                                                                                                    GASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
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                                                                                                                                                                                                                                                                                                                                                                                                                                      84.3%; Score 2589; DB 14;
85.1%; Pred. No. 1.4e-155;
tive 29; Mismatches 52;
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APPLICANT: BIGGEN, INC.
TITLE OF INVENTION: ANTIBODIES TO VLA-1
FILE REFERENCE: A101 PCT
CURRENT APPLICATION UNMBER: US/10/474,832
CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/233,794
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR FILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 70
SOPTWARE: PATENTIN VET. 2.1
SEQ ID NO 4
LENGTH: 447
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US-10-474-832-4
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Best Local S
Matches 426
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                                                                                                                                 SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                               LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
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                                                 TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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Pred. No. 4.8e-134;
4; Mismatches 17;
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GENERAL INFORMATION:
APPLICANT: BIOCEN, INC.
TITLE OF INVENTION: ANTIBODIES TO VLA-1
FILE REFERENCE: A101 PCT
CURRENT FILING DATE: 2003-10-14
CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-04-04
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR FILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 70
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.1
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LENGTH: 447
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Best Local Similarity 94.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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               QGNVFSCSVMHEALHNHYTQKSLSLSPG 448
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                                                                          TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                       TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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QGNVFSCSVMHEALHNHYTQKSLSLSPG
                                                           TKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWO
                                                                                                                      TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
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Pred. No. 1.1e-133;
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RESULT 8
US-10-412-406-32
; Sequence 32, Application US/10412406
; Publication No. US20040058394A1

Length 4852;

1; Gaps

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; LENGTH: 663
; TYPE: PRT
; ORGANISM: Homo S
US-10-412-406-32
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APPLICANT: GARBER, Ellen
APPLICANT: LYNE, Paul
APPLICANT: SALDHANA, JOSE W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINALOCCE
CURRENT APPLICATION NUMBER: US/10/412,406
CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: 60/240,285
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/275,289
PRIOR PILING DATE: 2001-03-13
PRIOR FILING DATE: 2001-03-13
PRIOR PILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/299,987
PRIOR APPLICATION NUMBER: FOT/US01/32140
PRIOR APPLICATION NUMBER: POT/US01/32140
PRIOR PILING DATE: 2001-10-12
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US-10-412-406-33
; Sequence 33, Application US/10412406
; Publication No. US20040058394A1
; GENERAL INFORMATION:
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Best Local Similarity
Matches 422; Conserv
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APPLICANT: BIOGEN, INC.
APPLICANT: GARBER, Ellen
APPLICANT: LYNE, Paul
APPLICANT: SALDHANA, Jose W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINA100CN
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SOFTWARE: FastSEQ for Windows Version
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                                                           TITLE OF INVENTION: ANTIBODIES TO VLA-1
FILE REFERENCE: A101 PCT
CURRENT APPLICATION NUMBER: US/10/474,832
CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR FILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 70
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Publication No. US20040081651A1
GENERAL INFORMATION:
APPLICANT: BIOGEN, INC.
                     SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows
SEQ ID NO 33
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Best Local Similarity
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ORGANISM: Homo
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Pred. No. 2.5e
8; Mismatches
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FILE REFERENCE: 7430*163
CURRENT APPLICATION NUMBER: US/10/320,231A
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: US 60/342,174
PRIOR FILING DATE: 2001-12-17
NUMBER OF SEQ ID NOS: 85
SOFTWARE: Patentin version 3.2
SEQ ID NO 79
LENGTH: 445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT; ORGANISM: Artificial Sequence; PEAGUURE: OTHER INFORMATION: Description; OTHER INFORMATION: polypeptide US-10-474-832-6
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US-10-320-231A-79
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Best Local S
Matches 424
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            Query Match
Best Local Similarity
Matches 421; Conserv
                                                            FEATURE:
OTHER INFORMATION:
-10-320-231A-79
                                                                                                                                                                                                                                         APPLICANT: Neben, Steven
APPLICANT: Takeuchi, Toshihiko
APPLICANT: Takeuchi, Toshihiko
APPLICANT: Tomkinson, Adrian
TITLE OF INVENTION: Antibody Inhibiting Stem Cell Factor Activity And Use
TITLE OF INVENTION: Treatment Of Asthma
                                                                                                     ORGANISM: Artificial
                                                                                                                  TYPE: PRT
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              Conservative
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                                                                           synthetic sequence
                      72.8%;
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            Score 2234.5; D
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RESULT 12
US-10-150-475A-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 6
LENGTH: 444
TYPE: PRT
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Best Local S
Matches 421
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Publication No. US20030103985A1

GENERAL INFORMATION:
APPLICANT: Adolf, G. et al.
TITLE OF INVENTION: Cytotoxic CD44 Antibody
FILE REFERENCE: 1/1211
CURRENT APPLICATION NUMBER: US/10/150,475A
CURRENT FILING DATE: 2002-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/307,451
PRIOR FILING DATE: 2001-07-24
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver: 2.1
                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Humanised OTHER INFORMATION: Murine Antibody BIWA 4 Heavy Chain SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
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                                                                                                             STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPBPVTVSWNSGALTSGVHTFPAVLQSSG
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                                     STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
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APPLICANT: Adolf, Gunther
APPLICANT: Baumann, Michael
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Compositions and methods for treating cancer using
TITLE OF INVENTION: Compositions and methody Immunoconjugates
FILE REFERENCE: 1/1414
CURRENT APPLICATION NUMBER: US/10/704,522
CURRENT FILING DATE: 2003-11-07
PRIOR APPLICATION NUMBER: US 60/429,516
PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: US 60/429,516
PRIOR FILING DATE: 2002-11-08
NUMBER OF SEQ ID NOS: 9
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 6
FIRMSTH: 444
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; OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Heavy
US-10-704-522-6
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US-10-704-522-6
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Best Local Similarity
Matches 421; Conserv
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TYPE: PRT
ORGANISM: Artificial Sequence
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TYRVVSVLTVLHQDWLNGKEYKCKVSNKAL.PAPIEKTISKAKGQPREPQVYTL.PPSRDEL
                                                                 SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                         SVFLFPPKPKDTLMI SRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRBEQYNS
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APPLICANT: Adolf, Guenther

APPLICANT: Baum, Anke

APPLICANT: Heider, Karl-Heinz

ITILE OF INVENTION: Compositions and Methods for Treating Cancer using

ITILE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and

ITILE OF INVENTION: Chemotherapeutic Agents

FILE REFERENCE: 1/1383

CURRENT APPLICATION NUMBER: US/10/645,215

CURRENT FILING DATE: 2003-08-21

PRIOR FILING DATE: August 21, 2002

PRIOR APPLICATION NUMBER: US 60/405,956

PRIOR APPLICATION NUMBER: US 60/405,956

PRIOR FILING DATE: August 26, 2002

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin Ver. 2.1
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US-10-645-215-6
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Publication No. US20040126379A1
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TYPE: PRT
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QGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ 420
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CURRENT APPLICATION NUMBER: US/09/736,371B
CURRENT FILING DATE: 2002-04-25
PRIOR APPLICATION NUMBER: 9815909.8
PRIOR FILING DATE: 1998-07-21
PRIOR APPLICATION NUMBER: PCT/GB99/02380
PRIOR APPLICATION NUMBER: PCT/GB99/02380
PRIOR FILING DATE: 1999-07-21
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 21
LENGTH: 449
TYDE: PRT
ORGANISM: Homo sapiens
US-09-736-371B-21
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US-09-736-371B-21
; Sequence 21, Application US/09736371B
; Patent No. US20020131968A1
; GENERAL INFORMATION:
Search completed: December 23, 2004, 19:35:46 Job time: 134.396 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Waldmann, Herman
APPLICANT: Frewin, Mark
TITLE OF INVENTION: AGLYCOSYLATED ANTIBODIES
FILE REFERENCE: Waldmann
                                                             421
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.

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	24.4	25.8	26.4	26.6	26.7	27.9	28.2	34.7	35.2	35.6	35.9	36.2	36.4	36.5	36.6	36.7
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	Ig gamma chain - m	Ig gamma-1 heavy c	Ig heavy chain V-I	Ig Y heavy chain (Ig heavy chain (DO	Ig heavy chain VHI	Ig heavy chain pre	Ig gamma 4 chain c	Ig gamma-2b chain	Ig gamma-2 chain C	Ig epsilon chain C		Ig gamma-2c chain		gamma-2a	Ig gamma-2a chain

Ig gamma-1 chain C region - human (\$Species: Homo sapiens (man) (\$Species: Homo sapiens (man) (\$C;Species: Homo sapiens (man) (\$C;Accession: A93433; S36861; \$\overline{S33887}; B90563; A90564; B91\overline{G8}; A91723; A02146 (\$Accession: A93433; S36861; \$\overline{S33887}; B90563; A90564; B91\overline{G8}; A91723; A02146 (\$R;Ellison, J.W.; Berson, B.J.; Hood, L.E. Nucleic Acids Res. 10, 4071-4079, 1982 (\$Accession: Ap3433; MUID:82274238; PMID:6287432 R;Harris, L.J. submitted to the EMBL Data Library, October 1992 A;Reference number: S33904 A;Cross-references: EMBL:Z17370 R;Takahashi, N.; Ueda, S.; Obata, M.; Nikaido, T.; Nakai, S.; Honjo, T. Cell 29, 671-679, 1674 A;Title: Structure of human immunoglobulin gamma genes: implications for evolution A;Reference number: S33887; MUID:83001943; PMID:6811139 A;Accession: S33887; A;Molecule type: DNA A;Residues: 1-330 <ELL> A;Residues: 1-350 <ELL> A;Cross-references: UNIPROT:P01857; EMBL:Z17370 A;Note: this sequence has the GIm(17) allotypic marker, 97-Lys, and the GIm(1) markers, A;Note: Lys-330 is removed after translation A;Note: Lys-330 is removed after translation A;Molecule type: DNA A;Residues: 2-330 <HAR> A;Accession: S36861 Accession: A93433;

A;Molecule type: protein
A;Molecule type: protein
A;Residues: 136-154,'Q',156-165,'Q',167-176,'Q',178-194,'N',196-197,'D',199-238,'E',240
A;Note: this sequence has the G1m(non-1) markers, 239-Glu and 241-Met
A;Ponstingl, H.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 357, 1571-1604, 1976
A;Title: Die Primaerstruktur eines monoklonalen IgG1-Immunglobulins (Myelomprotein Nie)

,240,

sequ.

A;Accession: A90564

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A;Molecule type: protein
A;Residues: 1-96, 'R', 98-135 <CUN>
A;Residues: 1-96, 'R', 98-135 <CUN>
A;Residues: 1-96, 'R', 98-135 <CUN>
A;Note: this sequence has the Glm(3) marker, 97-Arg
A;Note: this sequence has the Glm(3) marker, 07-Arg
A;Rutishauser, U; Cunningham, B.A.; Bennett, C.; Konigsberg, W.H.; Edelman, G.M.
Biochemistry 9, 3171-3181, 1970
Biochemistry 9, 3171-3181, 1970
A;Title: The covalent structure of a human gammaG-immunoglobulin. VIII. Amino acid
A;Reference number: A90564; MUID:71064025; PMID:5530842

A;Molecule type: DNA
A;Residues: 88-113;235-330 <TAK>
A;Residues: 88-113;235-330 <TAK>
A;Cross-references: EMBL:Z17370
A;Cross-references: EMBL:Z17370
R;Cunningham, B.A.; Rutishauser, U.; Gall, W.B.; Gottlieb, P.D.; Waxdal, M.J.; Edelman, Biochemistry 9, 3161-3170, 1970
Biochemistry 9, 3161-3170, 1970
A;Title: The covalent structure of a human gammaG-immunoglobulin. VII. Amino acid seque A;Reference number: A90563; MUID:71064024; PMID:5489771
A;Contents: myeloma protein Eu
A;Accession: B90563

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igen Primaerstruktur:

A; Reference number: Ap1668; MUID:77070269; PMID:826475

A; Contents: myeloma protein M;
A; Accession: Bp1668

A; A) Concinents: myeloma protein M;
A; Residues: 1-34, 'Q', 36-96, 'K', 98-115, 'Q', 117-197, 'D', 199-238, 'D', 240, 'L', 242-268, 'E', 27
A; Noce: this sequence has the Gim(17) and Gim(1) markers
R; Schmidt, W. E.; Jung, H. D.; Palm, W.; Hilschmann, N.
Hoppe-Seyler's Z. Ephysiol. Chem. 364, 713-747, 1983

A; Title: Die Primaerstruktur des kristallisierbaren monoklonalen Immunglobulins IgG1 KOI
A; Reference mumber: A91723; MUID:82289131; PMII:6848994
A; Contents: myeloma protein KO; disulfide bonds
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Residues: 1-96, 'R', 98-197, 'D', 199-238, 'E', 240, 'M', 242-266, 'D', 268-271, 'D', 273-330 <SCH
A; Title: The covalent structure of a human gammaG-immunoglobulin AgammaG-immunoglobulin herestotetermer subunit consists of two identical light (kaphyondes) and the consists
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;243-310/Domain: immunoglobulin homology <IM3>
;243-310/Domain: immunoglobulin homology <IM3>
;27-83,144-204,250-308/Disulfide bonds: #status experimental
;103/Disulfide bonds: interchain (to light chain) #status experimental
;109,112/Disulfide bonds: interchain (to heavy chain) #status experimental
;180/Binding site: carbohydrate (Asn) (covalent) #status experimental
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Best Local Similarity
Matches 330; Conserv
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                                                                                                                                        LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
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                                                  QQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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Pred. No. 5.9e-92;
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A23511

Ig gamma-3 chain C region (allotype G3m(b)) - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 28-Dec-1987 #sequence_revision 28-Dec-1987 #text_change
C;Accession: A23511
R;Huck, S:, Fort, P:, Crawford, D.H.; Lefranc, M.P.; Lefranc, G.
Nucleic Acids Res. 14, 1779-1789, 1986
A;Title: Sequence of a human immunoglobulin gamma 3 heavy chain
A;Reference number: A23511; MUID:86148507; PMID:3081877
A;Accession: A2351
A;Molecule type: DNA
A;Residues: 1-377 <HUC>
A;Cross-references: GB:X03604; GB:M12958; NID:g33070; PIDN:CAA27
C;Genetics:
C;Genetics:
A;Title: A human immunoglobulin IGHG3
A;Reference number: A60764; MUID:9000
A;Accession: A60764
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-377 <a href="https://doi.org/10.1016/nd/479">https://doi.org/10.1016/nd/479</a>
C;Superfamily: immunoglobulin C regic
C;Keywords: immunoglobulin
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A;Cross-references: GDB:119339; OMIM:147120
A;Map position: 14q32.33-14q32.33
A;Introns: 98/3; 115/3; 130/3; 145/3; 160/3; 270/3
A;Superfamily: immunoglobulin
C;Keywords: immunoglobulin
C;Keywords: immunoglobulin
bomology <IMM>
                                                                                                                                                                     R;Huck, S.; Lefranc, G.; Lefranc, Immunogenetics 30, 250-257, 1989
                                                                                                                                                                                                   Ig gamma-3 chain C region, form LAT - h
C;Species: Homo sapiens (man)
C;Date: 14-May-1993 #sequence_revision
C;Accession: A60764
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C;Species: Homo sapiens (man)
C;Date: 30-Apr-1981 #sequence revision 13-Jun-1983 #text_change 09-Jul-201
C;Date: 30-Apr-1981 #sequence revision 13-Jun-1983 #text_change 09-Jul-201
C;Accession: A93906, A92809; A92809; A93132; A02148
R;Ellison, J; Hood, L
Proc. Natl. Acad. Sci. U.S.A. 79, 1984-1988, 1982
A;Title: Linkage and sequence homology of two human immunoglobulin gamma l
A;Reference number: A93906; MUID:82197621; PMID:6804948
A;Accession: A93906
A;Accession
                                                                                        A,Title: The amino acid sequences of the three heavy chain A;Reference number: A90752; MUID:80001357; PMID:113060 A;Contents: myeloma protein Zie A;Accession: A90752 A;Molecule type: protein A;Residues: 1-24,'E', 26-57,'EV', 60-85;132-171,'ZZZ',175,'B'A;Hote: this sequence has since been revised R;Hofmann, T.; Parr, D.M.
Mol. Immunol. 16, 923-925, 1979
                                                                                                                                                                                                                                                                                                                                                       A;Title: The primary structure of a human IgG2 heavy chain: A;Reference number: A92809; MUID:81007873; PMID:6774012
A;Contents: myeloma protein Til
A;Accession: A92809
A;Molecule type: protein
A;Residues: 1-19,'Q','21-57,'Z','59,'A','61-193,'D','195-325 <WA;Residues: 1-19,'Q','21-57,'Z', 59,'A','61-193,'D','195-325 <WA;Note: Trp-156 is at or near the complement-binding site R;Connell, G.E.; Parr, D.M.; Hofmann, T.
Can. J. Biochem. 57, 758-767, 1979
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                                 the amino acid sequence of residues 381-391 A93132; MUID:80114419; PMID:118920
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Pred. No. 3.9
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                                                                                                                                                                                    ,175,'B',177-193,'D',195-196,'Q',198
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                                                                   human
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RESULT 5
G4HU
Ig gamma-4 chain C region - human
C;Species: Homo sapiens (man)
C;Date: 02-Apr-1982 #sequence revision 02-
C;Accession: A90933; A90249; A02150
C;Accession, J.; Buxbaum, J.; Hood, L.
DNA 1, 11-18, 1981
A;Title: Nucleotide sequence of a human im
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A;Cross-references: GDB:119338; OMIM:147110
A;Map position: 14q32.33-14q32.33
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kapparament); In some cases, such as IgA and IgM, the subunits associate into 1 C;Superfamily: immunoglobulin C region; immunoglobulin homology C;Keywords: duplication; glycoprotein; heterotetramer; immunoglobulin F;20-85/Domain: immunoglobulin homology <IM1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biochem. J. 121, 217-225, 1971

A;Title: Disulphide bridges of the heavy chain of human: A;Reference number: A90253; MUID:72033500; PMID:4940472
A;Reference annotation; myeloma protein Sa, disulfide box R;Frangione, B.; Milstein, C.; Pink, J.R.L.
Mature 221, 145-148, 1969
A;Title: Structural studies of immunoglobulin G.
A;Reference number: A93157; MUID:69064124; PMID:5782707
A;Contents: annotation; Sa, disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: protein
A;Residues: 238-275 <HOF>
R;Hofmann, T.; Parr, D.M.
submitted to the Atlas, March
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F;239-306/Domain: immunoglobulin homology <IM3>
F;14/Disulfide bonds: interchain (to light chain) #status experimental
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A; Contents: annotation; Zie,
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                           QQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                           STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                                                                                                                                                                        PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQFN
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                                                                                             MTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDKSRW
                                                                                                                                                                                             STFRVVSVLTVVHQDWLNGKEYKCKVSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREE
QQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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Pred. No. 7.7e-83;
2; Mismatches 13;
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02-Apr-1982

#text_change 09-Jul-2004

immunoglobulin

C-gamma4

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A;Reference number: A90933; MUID:83157104; PMID:6299662
A;Accession: A90933
A;Molecule type: DNA
A;Residues: 1-327 < ELL>
A;Residues: 1-327 < ELL>
A;Cross-references: UNIFROT:P01861
A;Residues: 1, 33.47, 1970
A;Title: Human immunoglobulin sublclasses. Partial amino acid sequence of the constant reference number: A90249; MUID:70207560; PMID:4192699
A;Title: Human immunoglobulin sublclasses. Partial amino acid sequence of the constant reference number: A90249; MUID:70207560; PMID:4192699
A;Reference number: A90249; MUID:70207560; 
monocional antibody 13-1 heavy chain - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 04-Feb-1998 #sequence_revision 13-Mar-1998 #text_change C;Accession: PC4436 R;Akashi, S.; Kato, K.; Torizawa, T.; Dohmae, N.; Yamaguchi, H.; Biochem. Biophys. Res. Commun. 240, 566-572, 1997 A;Title: Structural characterization of mouse monoclonal antibod A;Reference number: JC5810; MUID:98063277; PMID:9398605 A;Molecule type: protein A;Residues: 1-444 <AKA>
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C;Comment: This catalytic antibody has peroxidase oxidase activity. C;Superfamily: immunoglobulin C region; immunoglobulin homology F;251-320/Domain: immunoglobulin homology < IMM> F;251-320/Domain: immunoglobulin homology < IMM> F;251bsulfide bonds: interchain (to 98) #status predicted F;99/Disulfide bonds: interchain (to 109) #status predicted
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SNWEAGNTFTCSVLHEGLHNHHTEKSLSHSPGK
                                                                                                                                                                             LGGPSVFLFPFKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREE
---SSVFIFPFKPKDVLTITLTFKVTCVVVDISKDDPEVQFSWFVDDVEVHTAQTQPREE
                                                                                                                                                                                                                                                                                                                      SASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
                 SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                          RDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDK
                                                                                                                    QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPS
                                                                                                                                                                                                                                            D-LYTLSSSVTVPSSTWPSETVTCNVAHPASSTKVDKKIVPRDCG---
                                                                                                                                                                                                                                                                         SGLYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPP--CPAPEL
                                                                                                                                                                                                                                                                                                                                                                    KYAESVRGRFTISRDDSKSSVYLQMNRLREEDTATYYCCRTP-WVYAMDCWGQGTSVIVS
                                                                                                                                                                                                                                                                                                                                                                                                YYSDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVS
                                                          KEQMAKDKVSLTCMITDFFFEDITVEWQWNGQFAENYKNTQFIMDTDGSYFVYSKLNVQK
                                                                                                                                                                                                                                                                                                       SAKTTPPSVYPLAPGSAAQTNSMVTLGCLVKGYFPEPVTVTWNSGSLSSGVHTFPAVLQS
                                                                                                                                                                                                                                                                                                                                                                                                                               EVOXVETGGGLVRPGNSLKLSCLTSGFTFSNYRMHWLRQPPGKRLEWIAVITVKSDNYGA
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Pred. No. 2.5e
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.5e-77:
444
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A;Cross-references: EMBL:X62916; NID:g439; PIDN:CAA44699.1; PIDR;Symons, D.B.A.; Clarkson, C.A.; Beale, D.
Mol. Immunol. 26, 841-850, 1989

A;Title: Structure of bovine immunoglobulin constant region hea A;Reference number: \$06610; MUID:90097956; PMID:2513487

A;Reference number: \$06610; MUID:90097956; PMID:2513487

A;Residues: 142-470 <SYM>
A;Residues: 142-470 <SYM>
A;Cross-references: EMBL:X16701

A;Note: the sequence was determined from the germline gene C;Genetics:

A;Gene: Ig CH gamma-1

A;Introns: 98/1; 111/1; 221/1

C;Superfamily: immunoglobulin C region; immunoglobulin homology C;Keywords: glycoprotein; heterotetramer; immunoglobulin; membr F;161-225/Domain: immunoglobulin homology <IMM> Ig heavy chain precursor (B/MT.4A.17.H5.A5) - bo N;Alternate names: Ig gamma-1 chain C region (cl C.5pecies: Bos primigenius taurus (cattle) C;Date: 06-Jan-1995 #sequence revision 06-Jan-19 C;Accession: S22080; S06610; A31303 R;Sanders, P.G. submitted to the EMBL Data Library, November 199 A;Reference number: S22080 A;Status: preliminary A; Molecule type: mRNA A; Residues: 1-470 < SAN> November 1991 06-Jan-1995 PIDN:CAA44699.1; PID:g440 (clone 8.10) #text_change heavy chain gamma

immunoglobulin homology
r; immunoglobulin; membr

membrane

protein

_ and

gamm

٩,

257

417

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RESULT 8
$37483
$37483
Ig gamma-2a chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23.
C;Accession: $37483
R;Ducancel, F.F.D.
submitted to the EMBL Data Library, February 1993
A;Reference number: $37483
A;Accession: $37483
A;Accession: $37483
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-469 <DUC>
A;Cross-references: EMBL:X70423; NID:9406252; PIDN:CAA49868.1; PID C;Superfamily: immunoglobulin C region; immunoglobulin F;276-345/Domain: immunoglobulin homology <IMM>
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al Similarity 59.1%;
267; Conservative 7;
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                                                                                                                                                                                                                                          EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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AKTTAPSVYPLAPVCGDTTGSSVTLGCLVKGYFPEPVTLTWNSGSLSSGVHTPPAVLQSD 199
                        ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEFVTVSWNSGALTSGVHTFPAVLQSS 179
                                                                                                  NENFKGKATLTVDTSSSTAYMOLSSLTSEDTÄVYFCARAMGATATLLDYWGOGTTLTVSS
                                                                                                                                   SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT-YYFDSWGQGTLLTVSS
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                                                                                                                                                                                                                                                                                                         Score 1440.5; DB Pred. No. 1.7e-73; Pred. No. 1.7e-73; Pred. No. 1.7e-73; Pred. No. 108
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Pred. No. 4.3e-76;
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C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Superfamily: immunoglobulin C region; immunoglobulin; pyroglutamic acid
C;Keywords: disulfide bond; glycoprotein; immunoglobulin; pyroglutamic acid
F;1-446/product: Ig gamma-2a chain #status experimental <MAT>
F;1-117/Domain: V-D-U region <CVD>
F;1-117/Domain: C region <CCHP>
F;118-214/Domain: C region <CCHP>
F;118-214/Domain: C1 region <CCH2>
F;215-230/Region: hinge
F;21-340/Domain: C2 region <CCH2>
F;360-427/Domain: C3 region <CCH2>
F;360-427/Domain: C3 region <CCH3>
F;360-427/Domain: immunoglobulin homology <IMM>
F;360-427/Domain: immunoglobulin homology <IMM>
F;11Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;22-96,144-199,261-321,367-425/Disulfide bonds: #status predicted
F;132/Disulfide bonds: interchain (to light chain) #status predicted
F;224,227,229/Disulfide bonds: interchain #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Reference number: (A;Accession: S40295)
A;Accession: S40296)
A;Molecule type: protopy A;Residues: 1-446 <KI
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C;Species: Mus musculus (house mouse)
C;Date: 07-Apr.1994 #sequence_revision
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Best Local S
Matches 267
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                                                 181 LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPP--CPAPELLG
                                                                                                                                                        121
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1-446 <KLE>
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                                                                                                                             STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                                                                                                                                                            QIQLQQSGPELVRPGASVKISCKASGYTFTDYYIHWVKQRPGEGLEWIGWIYPGSGNTKY
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LYTLSSSVTVTSSTWPSQSITCNVAHPASSTKVDKKIEPRG-PTIKPCPPCKCPAPNLLG
                                                                                                                                                                                                       NEKFKGKATLTVDTSSSTAYMQLSSLTSEDSAVYFCARG--GKFAMDYWGQGTSVTVSSA
                                                                                                                                                                                                                                                SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSSA
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                                                                                                     KTTAPSVYPLAPVCGDTTGSSVTLGCLVKGYFPEPVTLTWNSGSLSSGVHTFPAVLQSD-
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                                                                                                                                                                                                                                                                                                                                                                                                                 46.8%; Score 1437; DB 2; 59.3%; Pred. No. 2.5e-73; tive 70; Mismatches 107
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177

180 118

236 238 120

60 60

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RESULT 10

S31459

Ig gamma-1 chain - sheep (fragment)
C;Species: Ovis orientalis aries, Ovis ammon aries (domest C;Species: Ovis orientalis aries, Ovis ammon aries (tomest C;Species: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_c C;Accession: S31459
R;Patri, S:, Mau, F.
submitted to the EMBL Data Library, December 1992
A;Reference number: S31459
A;Accession: S31459
A;Accession: S31459
A;Accession: S1459
A;Accession: S1459
A;Accession: S1459
A;Accession: S1459
A;Cross-references: EMBL:X69797
C;Superfamily: immunoglobulin C region; immunoglobulin hom C;Reywords; immunoglobulin homology <IMM>
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                  YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                             YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQP--ENNYKTTPPVLDSDGSFFL
                                                                                                                                            TKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV
                                                                                                                                                                                                                                                      TFPAILQSSGLYSLSSVVTVPASTSGAQTFICNVAHPASSTKVDKRVEPGCPDPCKHC-R
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                                                                                                                          TKPREEQFNSTFRVVSALPIQHQDWTGGKEFKCKVHNEALPAPIVRTISRTKGQAREPQV
                                                                                                                                                                                         CPPPELPGGPSVFIFPPKPKDTLTISGTPEVTCVVVDVGQDDPEVQFSWFVDNVEVRTAR
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                                                                                                                                                                                                                                                                                                                                                                                                                -TYYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYY-----FDSWG
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                                                             YVLAPPQEELSKSTLSVTCLVTGFYPDYIAVEWQKNGQPESEDKYGTTTSQLDADGSYFL
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RESULT

Ig gamma-2b chain - mouse (house mouse)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 31-Mar-1980 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004
C;Accession: S25057; A02157; A26235; A26232; A26233; A53598
R;Fischer, R.; Voss, A.; Niersbach, M.; Munziker, W.; Hirsch, H.J.; Kreuzal submitted to the EMBL Data Library, July 1992
submitted to the EMBL Data Library, July 1992
A;Description: Production of a Tobacco mosaic virus (TMV) inactivating neota; A;Accession: S25057
A;Accession: S25057
A;Status: preliminary

H.J.; Kreuzaler,

neotop specific

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Ig heavy chain V region precursor - human
(Species: Homo sapiens (man)
(Species: Homo sapiens (man)
(C)Species: Homo sapiens (man)
(C)Bate: 19-Mar-1997 ##equence_revision 19-Mar-1997 #text_change 01-Dec-2000
(C)Accession: S69339; 872664
R;Khamilchi, A.A.; Aucouturier, P.; Preud'homme, J.L.; Cogne, M.

Rur. J. Biochem. 229, 54-60, 1995
A;Title: Structure of abnormal heavy chains in human heavy-chain-deposition A,Reference number: 869339; MUID:95262687; PMID:7744049
A;Reference number: 869339; MUID:95262687; PMID:7744049
A;Status: preliminary
A;Status: preliminary
A;Status: 1-374 <KHA-
A;Residuse: 1-3
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A;Accession: 872664
A;Accession: preliminary
A;Molecule type: mRNA
A;Residues: 1-140,'C','142-374 <KH2>
A;Cross-references: EMBL:X81695
C;Superfamily: immunoglobulin C region; immunoglobulin homology
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Best Local Similarity
Matches 282; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RYSPSLRTRLTITKDTSKNQVVLTWINVDPADTATYYCGYSVEGYGQGYRFHSWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QITLKESGPTLVKPTQTLTLTCTFSGFSLSKSGVGVGWIRQPPGQALEWLALI-FWDDDK
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                       KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                      SREEMTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVD
                                                                                                                                                                          SRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
                                                                                                                                                                                                                                                                         EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
                                                                                                                                                                                                                                                                                                                EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
                                                                                                                                                                                                                                                                                                                                                                                                                LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                46.5%; ilarity 62.1%; Conservative 2
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Pred. No. 5.7e-73;
3; Mismatches 45
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A;Molecule type: DNA
A;Residues: 138-161,'L',163-189,'FP',193-300,'R',302-331,'A',333-437,'DI',440-474 <OLL>
A;Cross-references: GB:J00461
R;Kim, H: Yamaguchi, Y: Masuda, K: Matsunaga, C: Yamamoto, K: Irimura, T:; Takahash
J. Biol. Chem. 269, 12345-12350, 1994
A;Title: O-glycosylation in hinge region of mouse immunoglobulin G2b.
A;Reference number: A53598
A;Accession: A53598
A;Cession: A53598
A;Status: preliminary
A;Molecule type: protein
A;Residues: 234-251 <KIM>
C;Comment: The a allele sequence is shown.
C;Genetics:
A;Introns: 138/1; 236/1; 258/1; 368/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kaphain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: alternative splicing; duplication; glycoprotein; heterotetramer; immunoglobulin finance; immunoglobulin homology <IM1>
F;236-257/Region: hinge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A,Title: Structure of the constant and 3' untranslated regions A,Reference number: A26235; MUID:80081501; PMID:117548 A,Contents: MPC 11
A,Accession: A26225
A;Molecule type: mRNA
A,Residues: 138-172,'P',174-189,'FP',193-376,'T',378-474 <TUI>A;Note: Lys-474 is probably removed posttranslationally R,Tucker, P.W.; Marcu, K.B.; Newell, N.; Richards, J.; Blattner Science 206, 1303-1306, 1979
                                                                                                                                                                                                                                                                              F;236-257/Region: hinge
F;281-350/Domain: immunoglobulin homology <IM2>
F;281-350/Domain: immunoglobulin homology <IM2>
F;387-454/Domain: immunoglobulin homology <IM3>
F;387-454/Domain: imterchain (to light chain) #status predicted
F;152/Disulfide bonds: #status predicted
F;164-220,288-348,394-452/Disulfide bonds: #status predicted
F;247,250,253,256/Disulfide bonds: interchain (to heavy chain) #status
F;324/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Sequence of the cloned gene for the constant region A;Reference number: A26332; MUID:80081502; PMID:117549 A;Accession: A26332 A;Accession: A26332 A;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Note: the sequence was determined from the R;Tucker, P.W.; Marcu, K.B.; Slightom, J.L.; Science 206, 1299-1303, 1979
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A;Reference number: A26233; MUID:82173203;
A;Contents: b allele
A;Accession: A26233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 138-172,'P',1
R;Ollo, R.; Rougeon, F.
Nature 296, 761-763, 1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: DNA
A;Residues: 138-161,'L',163-189,'FP',193-474 <YAM>
A;Cross-references: GB:J00461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Complete nucleotide sequence of immunoglobulin A;Reference number: A02157; MUID:80120716; PMID:6766534 A;Contents: a allele A;Accession: A02157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:P01866; EMBL:X67210; NID:g54826; PIDN:CAA47649.1; R;Yamawaki-Kataoka, Y.; Kataoka, T.; Takahashi, N.; Obata, M.; Honjo, T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Residues: 138-172,'P',174-189,'FP',193-376,'T',378-474
                                                                                                                                                                                          Matches
                                                                                                                                                                                       260;
80
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                                                                                                                                                                                                                   Similarity
                                          SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTY---YFDSWGQGTLLTV 117
                                                                                                                                        EVOLVESGGDFVQFGGSLRVSCAASGFAFSHYAMSWVRQAFGKGLEWVAYISSGGSGTYY
                                                                                           EVQLQQSGPELVNPGASVKMSCKASGYTFITYVMHWVKQKPGQGLEWIGYINPNKDGTKF
                                                                                                                                                                                                              45.3%;
56.6%;
                                                                                                                                                                                          72;
                                                                                                                                                                                       Score 1390; DB 1;
Pred. No. 1.2e-70;
2; Mismatches 113;
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3; PMID:6803173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      germline of Blattner,
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                                                                                                                                 YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS
                                                                                                                                                                                                                                                     TKPRBEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV
                                                                                                                                                                                                                                                                                                                                                     CPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAK 290
                                                                                                                                                                                                                                                                                                                                                                                                                                   QSGLYTMSSSVTVPSSTWPSQTVTCSVAHPASSTTVDKKLEPSGPISTINPCPPCKECHK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SAAKTTPPSVYPLAPGCGDTTGSSVTSGCLVKGYFPESVTVTWNSGSLSSSVHTLSQALL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQ
KLNMKTSKWEKTDSFSCNVRHEGLKNYYLKKTISRSPGK
                                             KLTVDKSRWQQGNVFSCSVMHBALHNHYTQKSLSLSPGK 449
                                                                                                     YILPPPAEQLSRKDVSLTCLVVGFNPGDISVEWTSNGHTEENYKDTAPVLDSDGSYFIYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKT-HTCPP-----
                                                                                                                                                                                                                   TOTHREDYNSTIRVVSTLPIQHODWMSGKEFKCKVNNKDLPSPIERTISKIKGLVRAPOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 195
                                                                                                           435
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A;Cross-references: EMBL:X13188; NID:g51780; PIDN:CAA31580.1; PID:g51781
A;Note: this sequence was determined from the differentiated gene
C;Superfamily: immunoglobulin C region; immunoglobulin homology Ig gamma-2b chain precursor - mouse C;Species: Mus musculus (house mouse) C;Date: 30-Sep-1989 #sequence_revision F;1-19/Domain: signal sequence #status predicted <SIG>F;20-475/Product: Ig gamma-2b chain #status predicted F;159-223/Domain: immunoglobulin homology <IVM> A; Molecule type: mRNA A; Residues: 1-475 < DE1> A; Title: Expression in non-lymphoid cells of mouse record, Reference number: S01320; MUID:88329081; PMID:3138116 Ride Waele, P.; Feys, V.; van de Voorde, Eur. J. Biochem. 176, 287-295, 1988 C; Keywords: immunoglobulin A; Accession: S01321 Accession: Matches Query Match Best Local Similarity 257; 801321 EVOLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY Conservative 44.5%; Score 1367.5; DB Pred. No. 2.1e-69; 5; Mismatches 116 30-Sep-1989 A.; Molemans, 116; recombinant #text_change 2 ፕ : : Length Fiers, immunoglobulin 23-Jul-1999 9 Ξ directed

20 61 80

PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYT

APELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTK

292

198 232 258

318

APNIEGGPSVFIFPPNIKDVLMISLTPKVTCVVVDVSEDDPDVQISWFVNNVEVLTAQTQ

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKT-HTCPP-----CP

GLYTMSSSVTVPSSTWPSQTVTCSVAHPASSTTVDKKLEPSGPTSTINPCPPCKECHKCP

179

139

119

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319

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Iggamma 2a chain constant region - pig (fragment)
C;Species: Sue scrofa domestica (domestic pig)
C;Species: 1-Feb-1997 #sequence_revision 21-Feb-1997 #text_chang
C;Accession: 147159
R;Kaceskovics, I.; Sun, J.; Butler, J.E.
J. Immunol. 153, 3565-3573, 1994
A;Title: Five putative subclasses of swine IgG identified from A;Reference number: 147158; MUID:95015845; PMID:7930579
A;Accession: 147159
A;Accession: 147159
A;Katus: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-328 <KAC>
A;Cross-references: EMBL:U03779; NID:g433123; PIDN:AAA52217.1
C;Genetices:
A;Gene: IgG2a
C;Superfamily: immunoglobulin C region; immunoglobulin homolo
F;133-202/Domain: immunoglobulin homology <IMM'>
Ig gamma-1 chain C region - synthetic
C;Species: synthetic
A;Note: Homo sapiens (man) gene engineered and expressed in
C;Date: 06-Jan-1995 #sequence_revision 17-Mar-1997 #text_cha
C;Accession: S31866
R;Filpula, D.
submitted to the EMBL Data Library, February 1993
A;Description: Screeing method for protein-protein interacti
A;Reference number: S31866
A;Accession: S31866
A;Accession: S31866
A;Molecule type: mRNA
                                                                                                                                                                                             RESULT
S31866
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;Species: Sus scrofa domestica (domestic pig)
;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 21-Jan-2000
;Accession: I47159
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Best Local Similarity
Matches 231; Conserv
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; Pred. No. 9.4e
43; Mismatches
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A;Residues: 1-255 <FIL>
A;Cross-references: EMBL:X70421; NID:g33068; PIDN:CAA49866.1; PID:C;Keywords: immunoglobulin
F;1-22/Region: Bscherichia coli outer membrane protein A precursor
F;23-255/Region: human Ig gamma-1 chain C region
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PAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPE
                                                                                                                                    EDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKAL
                                                                                                                   EDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKAL
                                                                                                                                                                            TVAQADVESKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSH
                                                                                                                                                                                             TKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSH
                                                         PAPIEKTISKAKGOPREPOVYTLPPSRDELTKNOVSLTCLVKGFYPSDIAVEWESNGOPE
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                                                                                                                                                                                                                                                                                                                                         EMBL: X70421; NID: g33068; PIDN: CAA49866.1; PID: g33069
                                                                                                                                                                                                                                                   41.0%;
                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                   Score 1260; DB .
Pred. No. 1e-63;
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Search completed: Job time: 33.449 December secs 23, 2004, 19:06:04

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Result
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Maximum Match 100%
Listing first 45 su
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Perfect score:
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          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: uniprot_sprot:*
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1 EVQLVESGGDFVQPGGSLRV.....IVEFLNRWITFCQSIISTLT
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Gapop 10.0 , Gapext 0.5
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Q6PJA4
AAH18747
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AAH41037
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Q7z5w1 homo sapien
Q6p055 homo sapien
Aah65820 homo sapi
Bac85388 homo sapi
Bac85172 homo sapi
Aah64496 homo sapi
Bac85373 homo sapi
Bac85373 homo sapi
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Bac85232 homo sapien
Ofgmw7 homo sapien
Ofin78 homo sapien
Ofin78 homo sapien
Cae45781 homo sapien
Aah06402 homo sapien
Aah19046 homo sapien
Bac85444 homo sapien
Bac85444 homo sapien
Cae45774 homo sapien
Cae45774 homo sapien
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GRACES175 GRACES171 BACRES171 BACRES171 BACRES171 BACRES171 BACRES175 GRACES25 Q7Z7P5 GACRECASSES ALIGNMENTS ALIG	Qy 1 EVQLVESGGDFVQPGGSI : Db 20 EVQLVESGGLVKPGGSI	Query Match Best Local Similarity 92.9 Matches 420; Conservative	SEQUENCE FROM N.A. TISSUE=Spleen; Strausberg R.; Strausberg R.; Submitted (FEB-2002) to EMBL; BC024289; AAH24289 Hypothetical protein. SEQUENCE 471 AA; 5179	whiting M., Madan A., Whiting M., Madan A., Blakesley R.W., Touch Rodriguez A.C., Grimw Krzywinski M.I., Skal Jones S.J., Marra M.A. "Generation and initi and mouse cDNA sequen Proc. Natl. Acad. Sci [2]	SEQUENCE FROM N.A. TISSUE-Spleen; TISSUE-Spleen; MEDLINE-22388257; Pub Strausberg R.L., Pein Klausher R.D., Collin Altschul S.F., Zeeber Hopkins R.F., Jordan Diatchenko L., Marusi Stapleton M., Soares Brownstein M.J., Usdi Raha S.S., Loquellano Bosak S.A., McEwan P. Richards S., Worley K Villalon D.K., Muzny Faher, J. Holton P.	124289 PRELIMINARY; 124289; (TrEMBLrel. MAR-2004 (TrEMBLrel. MAR-2004 (TrEMBLrel. pothetical protein. no sapiens (Human). naryia; Metazoa; Chor mmalia; Eutheria; Prim	35 2107 68.6 481 36 2105.5 68.6 481 37 2105.5 68.6 481 38 2096 68.3 474 39 2084 67.9 472 40 2083.5 67.8 469 41 2082 67.8 470 42 2076 67.6 478 43 2075 67.6 478 44 2070 67.4 474 45 2069.5 67.4 474
Baceso Ba	RVSCAASGFAF	%; Score 2226.5; DB 2 %; Pred. No. 7.7e-137; 15; Mismatches 14;	he EMBL/GenBank/DDBJ database 1; MW; 388F7F4CF588660E CRC64;	Madan A., kodrigues Shevchenko Y., Bouff Shevchenko Y., Bouff een E.D., Dickson M.C utz J., Myers R.M., Blus D.E., Schnerch A. of more than 15,000 f 16899-16903 (2002).	E.A., Grouse L.H., Derge J.G. S., Wagner L., Shenmen C.M., Buetow K.H., Schaefer C.F., doore T., Max S.I., Wang J., Farmer A.A., Rubin G.M., H., Bonaldo M.F., Casavant T.L. Bonaldo M.F., Casavant T.L. J., Toshiyuki S., Carninci P., Toshiyuki S., Carninci P., Cokernan K.J., Malek J.A., Gu Hale S., Garcia A.M., Gay L. Sodergren E.J., Lu X., Gibe	PRT; 471 AA. Created) Last sequence update) Last annotation update) a; Craniata; Vertebrata; s; Catarrhini; Hominidae	175 773 171 125 225 236 336 017 017 012 013 013

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SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR--VKLGTY-YFDSWGQGTLLTV 117

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XX MEDLINE=22388257; PubMede=124777932;

XX MEDLINE=22388257; PubMede=124777932;

XX Klausner R.L., Feingold E.A., Grouse L.H., Derge J.G.,

XX Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

XX Altecchul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Altecchul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Altecchul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Altecchul S.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

XX Altecchul S.F., Zeeberg B., Buetow K.H., Casavant T.L., Scheet T.E.,

XX Altecchul S.F., Jordan H., Moore T., Casavant T.L., Scheet T.E.,

XX Altecchul S.F., Jordan H., Moore T., Casavant T.L., Scheet T.E.,

XX Altecchul S.F., Jordan H., Moore T., Marmason R.D., Mullahy S.J.,

XX Altecchul S.F., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

XX Altecchul S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

XX Altecchul S.F., Mockenn P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XX Altecchul S.F., Mocken P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XX Altecchul S., Mocken P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XX Altecchul S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XX Altecchul S.F., McKernan K.J., Malek J.A., Gunaratne P.H.,

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XX Altecchul S.F., McKernan K.J., Malek J.A., Gunaratne P.H.,

XX Altecchul S.F., McKernan 
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05-JUL-2004
05-JUL-2004
                           Strausberg R.;
Submitted (DEC-2001) to the
EMBL; BC018747; AAH18747.1;
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like
                                                                                                                                               SEQUENCE FROM N.
TISSUE=Primary E
                                                                                                                                                                                                                           Jones S.J., Marra M.A.;
"Generation and initial analysis
and mouse cDNA sequences.",
proc. Natl. Acad. Sci. U.S.A. 99:
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Mammalia; Eutheria;
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Homo sapiens (Human).
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       BC018747; AAH18747.1; -. Pro; IPR003599; Ig. Pro; IPR007110; Ig-like. Pro; IPR003597; Ig_c1.
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OC Euka
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InterPro; IPR003596; Ig_V.
Pfam; PF07654; C1-8et; 3.
Pfam; PF076047; 1g; 4.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGV; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 470 AA; 51715 MW; 7B49556
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AAH18747;
MEDLINE-22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.
                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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TISSUE=Primary B-Cells;
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Homo sapiens (Human).
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Hypothetical protein.
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Eukaryota, Metazoa; Chordata; Ci
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                                                                                                                                                                                                                                                                                                                                          Q6PI81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (DEC-2001) to the EMBL; BC018747; AAH18747.1; Hypothetical protein. SEQUENCE 470 AA; 51715 MW
  SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
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RX MEDLINE=22388257; PubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschall S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschanko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Foshiyuki S., Carninci P., Prange C.,

RA Staplèton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Wedin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Topes S. T. Marra M.A.
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"Generation and initial analysis
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99;
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10; Mismatches
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X MEDLINE-2388257; pubMed=12477932;
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X MEDLINE-2388257; pubMed=12477932;
X Manusher R.D., Collins F.S., Magner L., Scheefer C.F., Bhat N.K.,
X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
X Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
X Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
X Stapleton M.J., Usdin T.B., Toshiyuki S., Caraninci P., Prange C.J.,
X Stapleton M.J., Usdin T.B., Toshiyuki S., Caraninci P., Prange C.J.,
X Stapleton M.J., Usdin T.B., Toshiyuki S., Caraninci P., Prange C.J.,
X Stapleton M.J., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
X Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
X Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
X Halakes S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
X Wilalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
X Wilalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
X Wilalon D.K., Muzny D.M., Green E.D., Dickeon M.C.,
X Whiting M., Touchman J.W., Green E.D., Dickeon M.C.,
X Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
X Jones S. J. Marra M.A.
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02-MAR-2004 (TrEMBLrel.
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Proc. Natl. Acad. Sci. U.S.A.
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Mammalia; Eutheria;
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Pred. No. 6.9e-136;
0; Mismatches 20;
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Matches 416
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THE GERMAN HUMAN CDNA CONSORTIUM;
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Lauber J., Bahr A., Mewes H.W., Weil B., Amid C.,
Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ dat
EMBL; BX640947; CAE45972.1; -.
InterPro; IPR00359; IG.
InterPro; IPR00359; Ig.
InterPro; IPR003597; Ig-C1.
InterPro; IPR003597; Ig-C1.
InterPro; IPR003506; Ig-W.
Ffam; PF00647; IGR03506; Ig-W.
PFam; PF00047; IG; 4.
SWART; SW00409; IG; 2.
SWART; SW00409; IG; 2.
SWART; SW00406; IGW; 1.
PROSITE; PS00359; IG-IKE; 4.
PROSITE; PS00359; IG-IKE; 4.
PROSITE; PS00390; IG-MHC; UNKNOWN_2.
PROSITE; PS00390; IG-MHC; UNKNOWN_2.
SEQUENCE 475 AA; 52043 MW; B7EAE255A26F4B8E
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Q6MZQ6;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
                                                              AVLOSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPA
                                                                                   AVLQSSGLYSLSSVVTVPSSSLGTQTYI CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPA
                                                                                                                                         LVSVSAASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFP
                                                                                                                                                              LLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFF
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Pred. No. 7.4e
13; Mismatches
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Best Local S
Matches 416
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CAE45972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-Human esophagus tumor;
Lauber J. Bahr A., Mewes H.W.,
Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBI
EMBL; BX640947; CAE45972.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence up
02-MAR-2004 (TrEMBLrel. 27, Last annotation
Hypothetical protein DKFZp686G11190.
DKFZP686G11190.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae; Homo.
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  VDKSRWQQGNVFSCSVMHEGLHNHYTQKSLSLSPGK
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Pred. No. 7.4e-136;
3; Mismatches 20;
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Matches 409
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Fobo G., Han M., Wiemann S.;

Submitted (AUG-2003) to the EMBL/GenBank/DDBJ dat

EMBL; BX640853; CAR45920.1; -.

InterPro; IPR003599; Ig.

InterPro; IPR003599; Ig. 1ike.

InterPro; IPR003597; Ig. Cl.

R InterPro; IPR003597; Ig. MHC.

InterPro; IPR003595; Ig. W.

R Pfam; PP00654; C1-set; 3.

R Pfam; PP0047; Ig; 4.

SWART; SW00409; IG; 2.

SWART; SW00409; IG; 2.

SWART; SW00409; IG; 2.

SWART; SW00409; IG; 3.

R PROSITE; PS00835; IG LIKE; 4.

R PROSITE; PS00835; IG LIKE; 4.
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Q6MZV7;
Q6MZV7;
Q5-JUL-2004 (TrEMBLrel
Q5-JUL-2004 (TrEMBLrel
Q5-JUL-2004 (TrEMBLrel
Q5-JUL-2004 (TrEMBLrel
Q5-JUL-2004 (TrEMBLrel
Hypothetical protein D
Name=DKFZp686Cl1235;
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TISQUE-Human small intestine;
THE GERMAN HUMAN CDNA CONSORTIUM;
ROCCHET M., MEWES H.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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1 protein DKFZp686C11235.
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Primates;
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Catarrhini; Hominidae;
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Matches 409
                                                           BAC85232 PRELIMINARY;
BAC85232;
02-MAR-2004 (TrEMBLrel. 2
02-MAR-2004 (TrEMBLrel. 2
02-MAR-2004 (TrEMBLrel. 2
CDNA FLJ26265 fis, clone
Homo sapier Eukaryota;
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TISSUB=Human small intestine;
TISSUB=Human small intestine;
Bloecker H., Boecher M., Mewes H.W., Weil B.,
Bloecker H., Boecher M., Mewes H.W., Weil B.,
Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ
EMBL; BX640853; CAE45920.1; -.
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Mammalia; Eutheria; P
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Catarrhini;
  Craniata;
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MEDLINE=23.885.7; PubMed=12477932;
MEDLINE=23.885.7; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz
                                                                                                                                                                                                                                                                    Q6GMW7;
                                                                                                                                                                  Hypothetical protein.
Homo sapiens (Human).
Eukaryota, Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Ota T., Nakagawa S., Senoh A., Mizuguchi H., Inagaki H., St
Hata H., Nakagawa K., Mizuno S., Morinaga M., Kawamura M.,
Sugiyama T., Irie R., Otsuki T., Sato H., Nishikawa T., Su
Sugiyama T., Irie R., Otsuki T., Sugano S.;
"NEDO human CDNA sequencing project.";
Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AK129776; BAC85232.1; -.
SEQUENCE 472 AA; 51212 MW; O1BF215F99809164 CRC64;
                                                                                                                           SEQUENCE
                                                                                                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                        N.A.
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                                                                                                                                                                    Craniata; Vertebrata; Catarrhini; Hominidae
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No. 1.
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                                                                                                                                                                                 Euteleostomi;
                                                                                                                                                                     Homo.
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Query Match
Best Local S
Matches 415
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R EMBL; BC073782; AAH73782.1; -.

R InterPro; IPR003599; Ig.

R InterPro; IPR003597; Ig_c1.

R InterPro; IPR003597; Ig_c1.

R InterPro; IPR003597; Ig_wMC.

R InterPro; IPR003596; Ig_wMC.

R InterPro; IPR003596; Ig_w.

R Pfam; PF07654; C1-set; 3.

R Pfam; PF00047; Ig; 4.

R SMART; SM00409; IG; 2.

R SMART; SM00409; IG; 3.

R SMART; SM00406; IGv; 1.

R PROSITE; PS0035; IG_LIKE; 4.

R PROSITE; PS00290; IG_MIC; UNKNOWN_2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jones S.J., Marra M.A.;
Jones S.J., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. TISSUE=Spleen;
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Similarity 91.0%;
15; Conservative 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFP
                                                                                                                                                                                                                                                                                                                          PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
                                                                                                                                                                                                                                                                                                                                                                                                                           AVLQSSGLYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPA
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                                    VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                      PPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLT
                                                                                                                                                                                            REEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTL
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                                                                                                                                                                                                                                                                                                PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Score 2188.5; DB 2;
; Pred. No. 2.3e-134;
11; Mismatches 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; BCO72419; AAH72419; I;
InterPro; IPRO03599; Ig.
InterPro; IPRO03599; Ig.
InterPro; IPRO03597; Ig.cl.
InterPro; IPRO03597; Ig.cl.
InterPro; IPRO03597; Ig.mHC.
InterPro; IPRO03596; Ig.mHC.
INTERPO0407; Ig.g; 4.
SMART; SM00409; IG.; 2.
SMART; SM00409; IG.; 3.
SMART; SM00409; IG. INTERPOOSITE; PS50835; IG.LIKE; 4.
PROSITE; PS50835; IG.LIKE; 4.
PROSITE; PS50835; IG.LIKE; 4.
PROSITE; PS50835; IG.LIKE; 4.
PROSITE; PS50836; IG.LIKE; 4.
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Q6IN78;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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TISSUE=Peripheral Nervous System;
MEDLINE=22388257; PubMed=12477932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Peripheral Nervous System;
Strausberg R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.
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Eukaryota, Metazoa,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jones S.J., Marra M.A.; "Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=IGHG1;
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179
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SGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLG
                                                                   SASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
                                                                                                            SASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
                                                                                                                                                                                                         ADSVKGRFTISRDNSKNTLYLQMNSLRABDTAVYYCAR---GNYVVBAAPWGQGTLVTVS
                                                                                                                                                                                                                                                          EVQLVESGGGLIQPGGSLTLSCAASGLTVSSNYMHWVRQAPGKGLEWVSVLYIGGA-TYY
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Pred. No. 2.9e
11; Mismatches
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                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 414
                                                                                                                                                                                                                                                                                                                          Wambutt R., Heubhar D., Mewes H.W., Weil B., Amic Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ dat EMBL; Bx640627; CAE45781.1; -.
InterPro; IPR003599; Ig. Ilike.
InterPro; IPR003106; Ig_1ke.
InterPro; IPR003397; Ig_Cl.
InterPro; IPR003397; Ig_WHC.
InterPro; IPR003596; Ig_V.
Pfam; PF00647; Cl. set; 3.
pfam; PF00647; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGV; 1.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 472 AA; 51724 MW; 26CB340D0046D279 C
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686P15220.
Name=DKFZp686P15220;
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Eukaryota; Metazoa; (
Mammalia; Eutheria; E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Human rectum tumor;
THE GERMAN HUMAN CDNA CONSORTIUM;
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                                                                                                                             EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                          TVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSR
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   LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
                                                                                                           ADSVXGRETISRDNGKNSLYLQMNSLRAEDTALYYCAK-EIGAHNEYYYGMDVWGQGTTV
                                                                                                                                                                                  EVQLVESGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQAPGKGLEWVSGISWNSGSIAY
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                                                                                                                                                                                                                                                          Conservative
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Primates;
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                                                                                                                                                                                                                                                          13;
                                                                                                                                                                                                                                                      Score 2187; DB 2;
Pred. No. 2.9e-134;
3; Mismatches 21;
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Best Local Similarity
Matches 414; Conserv
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Submitted (AUG-2003) to the
EMBL; BX640627; CAE45781.1;
Hypothetical protein.
SEQUENCE 472 AA; 51724 MW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein DKFZP686P15220.
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02-mAR-2004 (TrEMBLrel. 27,
02-mAR-2004 (TrEMBLrel. 27,
02-mAR-2004 (TrEMBLrel. 27,
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SRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
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91.2%;
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae;
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Wambutt R., Heubner D., Mewes H.W.,
                                                                                                                                                                                                                                                                                                                                                                                                         EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                     EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
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EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
                                                                           LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
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el. 27, Last annotation updat
DKFZp686P15220
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Pred. No. 2.9e-134;
3; Mismatches 21;
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ANAHOGA OZ - MA
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RC TISSUE=Primary B-Cells;

RX MEDLINE=22388257; PubMed=12477932;

RX MEDLINE=22388257; PubMed=12477932;

RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Brownstein M.J., Uddin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., McZevan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Willalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

PA Jones S.J.
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Homo sapiens (Human).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation updat
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"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
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ALIGNMENTS

RESULT 1 AAB81988 ID AAB88 XX AAB8 XX AAB8 XX AAB8 XX AAB8 XX Gang XX Gang XX Gang XX Gang XX Canc XX Clai XX 30-SEP-1999; 99JP-00278291 06-APR-2000; 2000JP-00105088 29-SEP-2000; 2000WO-JP006774 05-APR-2001. WO200123432-A1. Synthetic. Ganglioside; GD3; complementarity determining region; CDR; antibody; Ganglioside GD3 03-JUL-2001 AAB81988 standard; protein; AAB81988; (KYOW) KYOWA HAKKO KOGYO KK (first entry) specific antibody related 108 B protein SEQ ID NO:

Query Match 100.0%; Score 568; DB 4; Best Local Similarity 100.0%; Pred. No. 1.2e-35; Matches 108; Conservative 0; Mismatches 0;

Length 108; Indels

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Gaps

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Sequence 108

A A

The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention

New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.

WPI; 2001-266143/27. Hanai N, Shitara K,

Nakamura

<u>,</u>7

Niwa

Claim 22; Page 172-173; 183pp; Japanese.

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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 128 AA;
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                        ABU11011 standard;
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                                                                                                                                                                                    RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLFWTFGQGTKVEIKR 108
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                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                        protein; 108
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                                                                                                                                                                                                                                                                                                                                                                Score 568; DB 4;
Pred. No. 1.4e-35;
Mismatches 0;
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RESULT 4
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drugs containing genetically-modified antibody against ganglioside GD3, its fragment, immunocompetent cell activators or/and antitumor agents i combination, applicable in treating malignant tumor like melanoma.
                                                                                                Ganglioside;
                                                                                                                          Ganglioside GD3 specific antibody related protein #3
                                                                                                                                                   03-JUL-2001
                                                                                                                                                                                                     AAB81994
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 8; Page 100; 121pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shitara K,
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                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                   DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAPKLLIFYSSNLHSGVPS
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                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                GD3;
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                                                                                                complementarity determining region; CDR; antibody;
                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                  98.9%;
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Pred. No. 3.4e-35;
0; Mismatches 1
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RESULT 5
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Best Local S
Matches 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a monoclonal antibody which can reac specifically with ganglioside GD3. The antibody and its derivatives useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and derivatives against g of e.g. tumors, with low activity in cancer.
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                                                                                                                                                                                                                                     30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ganglioside;
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06-APR-2000; 2000JP-00105088
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                                                                                                                                                                                             (KYOW)
human type complementation-determining region-transplanted derivatives against ganglioside GD3, useful in diagnosis ar
                                                                                             2001-266143/27.
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                                                                                                                                       Shitara K,
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Pred. No. 4.7e-35;
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Best Local 9
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                                              The present invention describes a monoclonal antibody which can reac specifically with ganglioside GD3. The antibody and its derivatives useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplifica of the invention
                                                                                                                                                                                                                                             New human type complementation-determining region-transplanted antibody derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ganglioside; cancer.
Sequence 128
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06-APR-2000; 2000JP-00105088
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                                                                                                                                                                                                  Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Hanai
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (KYOW ) KYOWA HAKKO KOGYO
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)B; AAF86895.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Shitara K,
                                                                                                                                                                                                  Page 153;
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98.1%;
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Pred. No. 1.3e-34;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                         Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ganglioside GD3 specific antibody related protein SEQ ID NO: 10.
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06-APR-2000; 2000JP-00105088
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Pred. No. 3.9e-34;
1; Mismatches 3
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                                                             Synthetic
                                                                                                                    Ganglioside;
                                                                                                                                                           Ganglioside GD3 specific antibody related protein #5.
                                                                                                                                                                                                    03-JUL-2001
                                                                                                                                                                                                                                           AAB81996;
                                                                                                                                                                                                                                                                               AAB81996 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 128 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   activity in cancer.
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06-APR-2000; 2000JP-00105088.
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                                                                                                                      GD3;
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95.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 545; DB 4;
Pred. No. 7.6e-34;
3; Mismatches 2
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                                                                                                                      region;
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                                                                                                                        antibody;
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RESULT 10
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Best Local S
Matches 103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
             WPI; 2001-266143/27.
N-PSDB; AAF86912.
                                                                                                                                                                                                                                         Ganglioside; GD3; complementarity
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06-APR-2000; 2000JP-00105088.
                                                                                             06-APR-2000;
                                                                                                        30-SEP-1999;
                                                                                                                               29-SEP-2000; 2000WO-JP006774
                                                                                                                                                        05-APR-2001
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                                                                                                                                                                                                                                                                Ganglioside
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DB; AAF86904.
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                                                                       куома накко кобуо кк
                                                                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           128 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                              DIOMTOSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPGKAPKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                 RESGGGSGTDYTLTISNLQPEDFATYYCHQYSKLPWTFGQGTKVEIKR
                                               Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 159-160; 183pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                               GD3 specific antibody related protein
                                                                                            99JP-00278291
2000JP-00105088
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                                                                                                                                                                                                                                                                                                                                    protein; 128
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                                                Nakamura
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Pred. No. 7.6e-34;
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                                                                                                                                                                                                                                         antibody;
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RESULT 11
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Best Local 9
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                                                                                                         New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
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                                                                                                                                                                                                  Hanai
                                                                                                                                                                                                                                           30-SEP-1999;
06-APR-2000;
                                                                                                                                                                N-PSDB;
                                                                                                                                                                            WPI;
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                                                                                                                                                                                                                                                                                                                                                                    cancer
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                                                                                                                                                                                                                                                                                                                                              Synthetic
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DB; AAF86892.
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                                                                                                                                                                                                 Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page
                                                                                                                                                                                                                                                                                                                                                                                                   GD3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                           99JP-00278291
2000JP-00105088
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                                                                                                                                                                                                                                                                                                                                                                              GD3;
                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                   specific antibody related protein
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                                                                                                                                                                                                                                                                                                                                                                             complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
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96.3%;
                                                                                                                                                                                                  Nakamura
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.1e-34;
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                                                                                                                                                                                                                                                                                                                                                                               antibody;
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The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention

exemplification

Example 1; Page 150-151; 183pp; Japanese

Sequence 128 AA;

Length 128

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AABB1990
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Best Local S
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Best Local S
Matches 101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ganglioside GD3 specific antibody related protein SEQ ID NO: 56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 11; Page 174-175; 183pp; Japanese.
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06-APR-2000; 2000JP-00105088
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity in cancer.
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5
                              61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
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                                                                                                                                 1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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                                                                                                                                                                                                              Conservative
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Pred. No. 7.3e-33;
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ismatches 5;
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                                                                                                                                                                                                                                                          Length 108,
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RESULT 13

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Location/Qualifiers

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Best Local S
Matches 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence
                        Mus
                                                                                                                                                                            AAB01628 standard;
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Drugs containing genetically-modified antibody against ganglioside GDJ, its fragment, immunocompetent cell activators or/and antitumor agents i combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ganglioside GD3;
                                                                                             Murine
                                                                                                                        07-DEC-2000
                                                                                                                                                  AAB01628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAR-2001; 2001JP-00097483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-2002; 2002WO-JP003170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modified ganglioside GD3 antibody associated protein
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                                                     humanised antibody
                                                                Mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200278739-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                         represents a protein associated with the anti- ganglioside GD3 antibody
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                                                                immunoglobulin;
                                                                                          immunoglobulin light chain variable region.
                                                                                                                                                                                                                                                                                                                                                            96;
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                                                                                                                                                                                                                                                                RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR
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                                                                                                                                                                            protein;
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                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                       90.5%;
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                                                                L chain; light chain; variable
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                                                                                                                                                                                                                                                                                                                                                         Score 514; DB 6;
Pred. No. 1.4e-31;
7; Mismatches 5
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Matches 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New humanized chimera antibody KM-871 useful for treating cancer, comprises variable region of mouse monoclonal antibody, reactive with ganglioside and human antibody constant region.
                                                                                                                                       Ganglioside; GD3; complementarity determining region; CDR; antibody; cancer.
                                                                                                                                                                              Ganglioside GD3 specific antibody related protein SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 128 AA;
 30-SEP-1999;
                           29-SEP-2000; 2000WO-JP006774.
                                                        05-APR-2001
                                                                                  WO200123432-A1
                                                                                                             Mus musculus
                                                                                                                                                                                                           03-JUL-2001
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18-SEP-1992;
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                                                                                                                                                                                                                                                             standard; protein; 128
                                                                                                                                                                                                                                                                                                                                  RFSGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                           (first entry
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92EP-00116026.
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/label= signal_peptide
21..128
/label= mature_immunoglobulin_light_chain_V_region
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                                                                                                                                                                                                                                                                                                                                                                                                New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                    Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                                                    activity in cancer.
                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 140; 183pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-266143/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hanai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (КУОМ ) КУОМА НАККО КОСУО КК.
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                                                           81
                                                                                  61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR
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                                                                                                                                1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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                                                       RFSGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                              DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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             2004,
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US-08-6458-516-12
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US-07-934-373C-17
US-08-146-206C-17
US-08-146-206C-17
US-09-705-9686-17
US-09-705-9728-17
US-09-705-992A-17
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Sequence 10, Appl Sequence 11, Appl Sequence 19, Appl Sequence 31, Appl Sequence 33, Appl Sequence 34, Appl Sequence 37, Appl Sequence 17, Appl Sequence 47, Appl Sequence 40, Appl Sequence 40, Appl Sequence 2, Appli Sequence 20, Appli Sequence 25, Appli
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Query Match 90.5%; Score 514; DB 4; Length 128; Best Local Similarity 88.9%; Pred. no. 1.2e-43; Matches 96; Conservative 7; Mismatches 5; Indels 0; Gaps Qy 1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVP Db 21 DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVP Qy 61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGGGTKVEIKR 108	TEMPLICATION NUMBER: US/09/225,322B TF FILING DATE: 1999-01-05 APPLICATION NUMBER: US 08/454,680 FILING DATE: 1995-05-31 APPLICATION NUMBER: US 08/408,133 FILING DATE: 1995-03-2 APPLICATION NUMBER: US 08/292,178 FILING DATE: 1994-08-17 APPLICATION NUMBER: US07/947,674 FILING DATE: 1992-09-17 APPLICATION NUMBER: JP 3-238375 FILING DATE: 1991-09-18 OF SEQ ID NOS: 19 OF SEQ ID NOS: 19 OF SEQ ID NOS: 19 RE: PATENTIN Ver. 2.0 NO 10 H: 128 PRT ILSM: Artificial Sequence RE: INFORMATION: Description of Artificial Sequence: INFORMATION: Description of Artificial Sequence:	ALIGNMENTS BSULT 1 S-09-225-322B-10 Sequence 10, Application US/09225322B Patent No. 6437098 GENERAL INFORMATION: APPLICANT: SHITARA, KENYA APPLICANT: HANAI, NOBUO APPLICANT: HANAI, NOBUO APPLICANT: HASEGAMA, MAMORU APPLICANT: HASEGAMA, MAMORU APPLICANT: MIYAJI, HIROMASA APPLICANT: KUWANIA, YOSHIHISA TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED	28 480 84.5 233 3 US-08-437-642B-25 29 480 84.5 233 4 US-09-705-686-25 30 480 84.5 233 4 US-09-705-686-25 31 480 84.5 233 5 US-09-705-392A-25 32 480 84.5 233 5 PCT-US93-07882-25 33 480 84.5 237 3 US-09-097-171A-10 35 480 84.5 237 3 US-09-097-172B-2 36 480 84.5 237 3 US-09-422-712B-2 36 480 84.5 237 3 US-09-407-756-2 37 480 84.5 237 3 US-09-407-756-2 38 480 84.5 237 3 US-09-407-756-2 39 490 84.5 237 3 US-09-40-166A-6 39 479 84.3 107 1 US-08-458-516-8 40 476 83.8 107 3 US-09-254-189-1 41 475 83.6 108 3 US-09-254-189-1 42 475 83.6 108 4 US-09-95-798-6 43 475 83.6 111 1 US-08-137-117D-67 45 475 83.6 111 1 US-08-137-117D-71
LLIFYSSNLHSGVPS 60 LLIFYSSNLHSGVPS 60 LLIFYSSNLHSGVPS 80 LLIFYSSNLHSGVPS 80 IKR 108 IKR 128	CDNA KM-641	CHIMERA ANTIBODY	Sequence 25, Appl Sequence 25, Appl Sequence 25, Appl Sequence 25, Appl Sequence 25, Appl Sequence 6, Appli Sequence 10, Appl Sequence 2, Appli Sequence 6, Appli Sequence 6, Appli Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3, Appli Sequence 67, Appli Sequence 67, Appli Sequence 67, Appli Sequence 67, Appli Sequence 67, Appli Sequence 67, Appli

RESULT 2 US-09-225-322B-19 ; Sequence 19, Application US/09225322B

INFORMATION:

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APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HANSGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITILE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR PILING DATE: 1995-08-17
PRIOR PILING DATE: 1995-08-17
PRIOR PILING DATE: 1995-08-17
PRIOR PILING DATE: 1995-08-17
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; APPLICANT: SHITARA, KENYA
; APPLICANT: HANAI, NOBUO
; APPLICANT: HASEGAWA, MAMORU
; APPLICANT: MIYAJI, HIROMASA
; APPLICANT: KUWANA, YOSHIHISA
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US-09-764-304-10
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Patent No. 6495666
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Best Local (
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NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 19
                                                                                                                                                      TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY FILE REPERENCE: 249-101

CURRENT PAPALICATION NUMBER: US/09/764,304

CURRENT FILING DATE: 2001-01-19

EARLIER APPLICATION NUMBER: 09/225,322

EARLIER FILING DATE: 1999-01-05

EARLIER APPLICATION NUMBER: US 08/454,680

EARLIER APPLICATION NUMBER: US 08/454,680

EARLIER APPLICATION NUMBER: US 08/408,133

EARLIER APPLICATION NUMBER: US 08/408,133

EARLIER APPLICATION NUMBER: US 08/292,178

EARLIER FILING DATE: 1995-03-21

EARLIER FILING DATE: 1994-08-17

EARLIER FILING DATE: 1994-08-17

EARLIER FILING DATE: 1994-08-17

EARLIER FILING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: UP 3-238375
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
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PRIOR FILING DATE: 1992-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: light chain OTHER INFORMATION: variable region
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Pred. No. 1.2e-43;
7; Mismatches 5
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US-09-764-304-19
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CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1994-08-17
EARLIER FILING DATE: 1992-09-17
                                                                                                                                                                                                                                                                                   SEQ ID NO 19
LENGTH: 128
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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LENGTH: 128
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Patent No. 6495666
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APPLICANT: HANAI, NOMO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                              OTHER INFORMATION: light chain OTHER INFORMATION: variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: cDNA KM-641
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                                                                                                                                                                       Local
96;
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                                                                                                                                                                       Similarity
                                                                                             DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
                                                                        DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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                                                                                                                                                                   90.5%;
                                                                                                                                                      7;
                                                                                                                                                  Score 514; DB 4;
Pred. No. 1.2e-43;
7; Mismatches 5
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Pred. No. 1.
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                                                                                                                                                                                         DB 4; Length 128,
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2e-43;
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RESULT 5 US-09-440-781-94

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CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 94
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Patent No. 606884
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                                                                                                                                                                                                                                                                                        COUNTY.
ZIP: 22314
ZIP: 22314
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: PC-DOS/MS-DOS
MEDIUM TYPE: PC-DOS/MS-DOS
MEDIUM TYPE: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SEINO, Ken-ichiro
APPLICANT: SEINO, Ken-ichiro
APPLICANT: KAYAGAKI, No. 6068841uhiko
APPLICANT: YAGITA, Hideo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 94, Application US/09440781 Patent No. 6632926
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                                                              TELEFAX: 703-684-1124
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: humanized antibody light chain variable domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: artificial sequence
                                                                                                                                                    CLASSIFICATION: 424
ATTORNBY/AGENT INFORMATION:
NAME: Bucca Ph.D., Daniel
REGISTRATION NUMBER: P-42,368
REFERENCE/DOCKET NUMBER: 50356-151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                   TELEPHONE: 703-518-5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 99 came
CITY: Alexandria
                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
            TYPE: amino acid
STRANDEDNESS: sin
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                                                                                                                                                                                                                                                                                                                                                                                                                       Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                      99 Canal Center Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OKUMURA, KO
NAKATA, MOLOMI
NENTION: THERAPEUTIC AGENT FOR HEPATITIS
                                            108 amino acids
                                                                                                   703-684-1124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McDermott, Will & Emery
linear
            single
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87.0%;
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Pred. No. 1.2e-42;
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Best Local Similarity
Thes 91; Conserve
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                                                                                                                                                                             ; MOLECULE TYPE: US-08-649-100-33
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                                                                                                                     Matches
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Patent No. 6
                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                             IELEFAX: (703) 205-8050
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                         NAME: MURPHY JR, GERALD M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 11:
TELECOMUNICATION INFORMATION:
TELEPHONE: (703) 205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: BIRCH, STEWART, KOLASCH AND BIRCH
STREET: PO BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                         TYPE: amino a STRANDEDNESS: TOPOLOGY: lin
                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: PO DOLL CTITY: FALLS CHURCH
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                    LENGTH:
 18
                    61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIK 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
                                                                                                                   92;
                                                                            1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RY: USA
22040-0747
                                                                                                                                                                                                                                     amino acid
                                                        DIQMTQSPSTLSASVGDRVTITCRASQDISNYLNWYQQKPGKAPKLLIYYTSRLHSGVPS
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RFSGSGSGTNYTLTISSLQPDDFATYFCQQGSTLPWTFGQGTKVEVK 127
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                                                                                                                                                                                                                                                    127 amino acids
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NAGATA, SHIGEKAZU
CO, MAN SUNG
                                                                                                                     Conservative
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                                                                                                                                                                                            peptide
                                                                                                                                                                                                                       single
                                                                                                                               85.7%;
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                                                                                                                Score 487; DB 3;
Pred. No. 5.6e-41;
6; Mismatches 9
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Pred. No. 1.9e-41;
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SING THE ANTI-FAS LIGAND ANTIBODY
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US-08-458-516-12
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Patent No.
                                                                                                                                                               Sequence 95, Applicate Patent No. 6632926 GENERAL INFORMATION:
NUMBER OF SEQ ID NOS:
SEQ ID NO 95
LENGTH: 110
TYPE: PRT
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Best Local (
                                                                          CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
                                                                                                            APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS FILE REFERENCE: P1469R1
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INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 214 amino acids
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APPLICANT: TBO, J. 1
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-37-3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower,
CITY: San Francisco
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 03-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single TOPOLOGY: linear
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o. 5777085
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                                                                                                                                                                                                                                                                                                                                                                                      1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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                                                                                                                                                                                                                                                                                                                                                                   DIQMTQTPSTLSASVGDRVTISCRASQDINNYLNWYQQKPGKAPKLLIYYTSTLHSGVPS
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                                                                                                                                                                                                     Application US/09440781
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83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 484; DB 1
Pred. No. 2e-40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 214;
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US-07-934-373C-17
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Matches
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Best Local
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APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
                                                                                                                                                                                                      TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                          FILING DATE: 21-Aug-1992
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US
FILING DATE: 15-UTUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 1-110
OTHER INFORMATION: humanized antibody light chain variable domain
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                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                               FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: IT NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WinPatin (Genentech)
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                                                                              Local
                                                                                                                                           TOPOLOGY:
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                                                                                                                                                        Amino Acid
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DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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The San Francisco
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Pred. No. 1.9e-40;
2; Mismatches 8
                                                                Score 480; DB 2;
Pred. No. 2.3e-40;
4; Mismatches 11;
                                                                                             Length 107;
                                                                 Indels
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RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIK 107

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US-08-437-642B-17
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US-08-652-558-2
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Patent No. 5861155
GENERAL INFORMATION:
                                                                                                                                        Patent No. 6054297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/652,558

FILING DATE: JUNE 6, 1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/IB94/00387

FILING DATE: NOVEMBER 21, 1994

ATTONNEY/AGENT INFORMATION:

NAME: YANKWICH, LEON R.

REGISTRATION NUMBER: 30,237

REFERENCE/DOCKET NUMBER: 30,237

REFERENCE/DOCKET NUMBER: 95,497-L

TELECOMMUNICATION INFORMATION:

TELEPHONE: 617-345-9100
GENERAL INFORMATION:
APPLICANT: Leanl J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSE: Genentech, Inc.
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette, 1.4
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION DATA:
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ADDRESSEE: BANNER & WITCOFF
STREET: 75 STATE STREET, 23F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: LIN, AUG
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                             Match 84.5%;
Local Similarity 86.8%;
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TOPOLOGY: linear
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                                                                                                                                                            Application US/08437642B
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Matches
                                                                                                                                                                                                          Sequence 17, Application US/08146206C Patent No. 6407213
GENERAL INFORMATION:
APPLICANT: Carter, Paul J.
APPLICANT: Presta, Leonard G.
APPLICANT: Method for Mal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/437,642B
FILING DATE: 09-May-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PILING DATE: 17-NOV-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc.
COMPUTER: IBM PC com.
OPERATING SYSTEM: PC
                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
                                                                                                                                        ADDRESSEE: GEAR
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STATE: C
COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
                                                                                                                                                                                                                                                                                                                                                                                                         61 RFSGSGSGTDYTLTISSLQPEDFATYYCQQGNTLPWTFGQGTKVEIK 107
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California
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WinPatin (Genentech)
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                         Genentech, Inc
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for Making Humanized Antibodies
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Pred. No. 2.3e-40;
4; Mismatches 11
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CURRENT APPLICATION DATA:

FILING DATE:

PPLICATION NUMBER: US/08/146,206C TLING DATE: 17-No. 6407213-1993

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RESULT 14
US-09-705-686-17
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GENERAL INFORMATION:
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Best Local (
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INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                  INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Lee, Wendy M.
REGISTRATION UNMBER: 40,378
REFERENCE/DOCKET NUMBER: PO:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 530
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 07/715272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M:
REGISTRATION NUMBER: 40,378
REGISTRATION INFORMATION:
TELEPHONE: 650/225-1994
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: Amino Acid TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb:
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/705,686
FILING DATE: 02-No. 6639055-2000
CLASSIFICATION: Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Carter, Paul J.

Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIK 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
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                TELEFAX: 650/952-9881
N FOR SEQ ID NO: 17:
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Pred. No. 2.3e-40;
                                                                                                                                                                                                                                                                                                                                                                                                           1.44 Mb floppy disk
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RESULT 15
US-09-705-392A-17
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                                                                                                                                                                                    ; SEQUENCE DESCRIPTION: SEQ ID NO: 17: US-09-705-392A-17
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GENERAL INFORMATION:
                                                                                                               Matches
                                                                                                                                 Best Local
                                                                                                                                                Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                  TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:

(UN100 DATA:

APPLICATION NUMBER: US/09/705,392A

FILING DATE: 02-00. 6719971-2002

CLASSIFICATION: «Unknown»

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206

FILING DATE: 17-NOV-193

APPLICATION NUMBER: 07/715272

FILING DATE: 11-UN-1991

ATTORNEY/ACENT INFORMATION:

NAME: Lee, Wendy M.

REGISTRATION NUMBER: 40,378

REFERENCE/DOCKET NUMBER: P0709P1D1 RE
TELEPHONE: 650/225-1994

TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94080
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Carter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIK 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 RFSGSGSGTDYTLTISSLQPEDFATYYCQQGNTLPWTFGQGTKVEIK 107
 61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIK 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 DIQMTQSPSSLSASVGDRVTITCRASQDIRNYLNWYQQKPGKAPKLLIYYTSRLESGVPS
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                                                                          1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
                                                                                                                                 Similarity
                             TYPE: Amino Acid TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOPTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: 1 DNA Way
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                  LENGTH: 107 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 107 amino
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                                                                                                               Conservative
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86.0%;
                                                                                                                                 84.5%; Score 480; DB 4; Length 107; 86.0%; Pred. No. 2.3e-40;
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                                                                                                               4; Mismatches
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Pred. No. 2.3e-40;
                                                                                                                                                                                                                                                                                                                                                                                  P0709P1D1 REVISED
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61 RFSGSGSGTDYTLTISSLQPEDFATYYCQQGNTLPWTFGQGTKVEIK 107

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Search completed: December 23, 2004, 19:08:00 Job time : 17.1328 secs

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                               Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                December 23, 2004, 19:04:51 ; Search time 24.5684 Seconds (without alignments) 1578.640 Million cell updates/sec
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1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
         GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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/Ggn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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/Ggn2_6/ptodata/2/pubpaa/US09B_PUBCOMB.pep:*
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/Ggn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
/Ggn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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    US-09-056-160B-103

US-10-234-671-101

US-09-056-160B-100

14 US-10-234-671-100

13 US-10-011-125-2

US-09-056-160B-15

US-10-234-671-15
                                                                                                                                                              US-09-764-304-10
US-09-764-304-19
4 US-10-265-713-10
4 US-10-265-713-19
4 US-10-166-626-19
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Sequence 10, Appl
Sequence 19, Appl
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Sequence 10, Appl
Sequence 10, Appl
Sequence 103, App
Sequence 101, App
Sequence 100, App
Sequence 100, App
Sequence 100, App
Sequence 20, Appl
Sequence 15, Appl
Sequence 15, Appl
                                                                                                                                                                                                                                                                                                                          Description
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ALIGNMENTS

US-09-764-304-10

Sequence 10, Application US/09764304
Patent No. US20020026036A1
GENERAL INFORMATION:

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GENERAL INFORMATION: GHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASGGAWA, MAMORU
APPLICANT: HASGGAWA, MAMORU
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
FULE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT APPLICATION NUMBER: US/02/25,322
EARLIER APPLICATION NUMBER: US/02/25,322
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-09-17
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SEQ ID NO 10
LENGTH: 128
TYPE: PRT
GRANUSM: Artificial Sequence
FEATURE:
FORMATION: CDNA KM-641
US-09-764-304-10
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US-09-764-304-19
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CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US 07/947,674
EARLIER FILING DATE: 1994-08-17
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-17
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Best Local S
Matches 96
              Sequence 10, Application US/10265713
Publication No. US20030095964A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HASEGAWA, MAMORU
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 19
LENGTH: 128
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: JP 3-238375
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 249-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: light chain OTHER INFORMATION: variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                      21 DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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KUWANA, YOSHIHISA
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                                                                                                                                                                                                                                                                                                                                                                                                                       Score 514; DB 9; Length 128; Pred. No. 7.1e-37;
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FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR PPLICATION NUMBER: US 08/454,680
PRIOR PPLICATION NUMBER: US 08/454,680
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR APPLICATION NUMBER: US007/947,674
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-265-713-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19, Application US/10265713 Publication No. US20030095964A1 GENERAL INFORMATION:
                                                                                                                           SEQ ID NO 19
LENGTH: 128
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SEQ ID NO 10
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TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR PILING DATE: 1999-01-05
PRIOR PILING DATE: 1999-01-05
PRIOR PILING DATE: 1995-05-31
PRIOR FILING DATE: 1995-05-31
PRIOR PILING DATE: 1995-05-31
PRIOR PILING DATE: 1995-03-21
PRIOR PILING DATE: 1995-03-21
PRIOR PILING DATE: 1995-08-17
                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: UP 3-238375
PRIOR FILING DATE: 1991-09-18
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                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 19
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                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                TYPE: PRT
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OTHER INFORMATION: Description of Artificial Sequence: light chain
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MIYAJI, HIROMASA
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RESULT 6
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Best Local S
Matches 96
                                                                          Sequence 19, Application US/10166626 Publication No. US20030166876A1
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SEQ ID NO 10
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                                                          GENERAL INFORMATION:
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PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMO
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PRIOR APPLICATION NUMBER: JP 3-238375
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CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR FILING DATE: 1999-01-05
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APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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Best Local Similarity
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PRIOR FILING DATE: 1999-01-05
PRIOR PRIOR ETLING DATE: 1995-05-31
PRIOR FILING DATE: 1995-05-31
PRIOR FILING DATE: 1995-05-31
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NUMBER OF SEQ ID NOS: 19
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYDE: 3.5 inch., 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                   PRIOR APPLICATION DATA: 60/054,856
                                                                                                               SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
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                                                                                                                                                                                                                                   ZIP:
                                                       APPLICATION NUMBER: US/09/056,160B FILING DATE: 06-Apr-1998 CLASSIFICATION: 424
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FILING DATE: 1995-03-21
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                                                                                                                                   WinPatin (Genentech)
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Presta, Leonard
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US-10-234-671-101
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SEQUENCE CHARACTERISTICS:
LENGTH: 110 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 101, Application US/10234671
Publication No. US20030190317A1
GENERAL INFORMATION:
APPLICANT: Baca, Manuel
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                                   INFORMATION FOR SEQ ID NO: 101:
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APPLICATION NUMBER: 09/056160

FILING DATE: 06-APR-1998

APPLICATION NUMBER: 60/126446

FILING DATE: 07-APR-1997

APPLICATION NUMBER: 60/054856

FILING DATE: 06-AUG-1997
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                                               ATTORNEY/AGENT INFORMATION:
NAME: Cui, Steven X.
REGISTRATION NUMBER: 44,637
REFERENCE/DOCKET NUMBER: P1093R2C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-8674
TELEPAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Windatin (Genencech)
CURRENT APPLICATION DATA:
               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/10/234,671 FILING DATE: 03-Sep-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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Lowman, Henry B.
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amino acids
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Pred. No. 2.5e-36;
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TOPOLOGY: Linear
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SEQUENCE DESCRIPTION: SEQ ID NO: 101:
US-10-234-671-101
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PRILING DATE: 06-Apr-1200

CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/054,856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: p1093R
TELECOMMUNICATION: NUMBER: 50/225-1896

TRI.EPHONE: 650/225-1896
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US-09-056-160B-100
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GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/056,160B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Baca, Manuel APPLICANT: Wells, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
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Local Similarity 88.0%;
les 95; Conservative
                                                                                                                                                   Local
                                                                                                                                                                                                                                 TYPE: Amino Acid
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                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                  LENGTH:
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84
                    61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
                                                              24 DIQUTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQXPGKAPKLLIYFTSSLHSGVPS
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                                                                                                                                            n 89.3%;
Similarity 88.0%;
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 RESGSGSGTDYTLTISSLQPEDFATYYCQQYSTVPWTFGQGTKVEIKR 131
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San Francisco
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Lowman, Henry B.
Chen, Yvonne M.
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                                                                                                                                                                                                                                                    237 amino acids
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                                                                                                                                 Conservative
                                                                                                                                                                                                                   Linear
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                                                                                                                                                                                                                                                                                                                                                       P1093R2
                                                                                                                                Score 507; DB 9;
Pred. No. 5.3e-36;
7; Mismatches 6
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                                                                                                                                                                  DB 9; Length 237;
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                                                                                                                                 Indels
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US-10-011-125-2
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Sequence 2, Application US/10011125
Publication No. US20020142388A1
GENERAL INFORMATION:
APPLICANT: Chen, Christina Yu-Ching
TITLE OF INVENTION: BACTERIAL HOST (
FILE REFERENCE: P1804R1
                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 100: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/054856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Cui, Steven X.
REGISTRATION NUMBER: 44,637
REFERENCE/DOCKET NUMBER: P1093R2C1
TELEPHONE: 650/25-8674
TELEPHONE: 650/25-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy diversely formatible computer: IBM PC compatible operating SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/234,671

FILING DATE: 03-Sep-2002

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Presta, Leonard G.
Lowman, Henry B.
Chen, Yoonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Baca, Manuel Wells, James A.
                                                                                                                                                                                                                                                                               24 DIQLTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPGKAPKLLIYFTSSLHSGVPS
                                                                                                                                                                                                         84 RFSGSGSGTDYTLTISSLOPEDFATYYCQQYSTVPWTFGQGTKVEIKR
                                                                                                                                                                                                                                             61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
                                                                                                                                                                                                                                                                                                                                                                         l Similarity
95; Conserv
                                                                                                                                                                                                                                                                                                                              1 DIOMTOSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 237 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/056160 FILING DATE: 06-APR-1998 APPLICATION NUMBER: 60/126446 FILING DATE: 07-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                        89.3%;
                                                                                                                                                                                                                                                                                                                                                                       Score 507; DB 14;
Pred. No. 5.3e-36;
7; Mismatches 6;
                       STRAINS
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                                                                                                                                                                                                                                                                                                                                                                         Indels
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US-09-056-160B-15
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US-09-056-160B-15
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/011,125
CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: US 60/256,162
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 10
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. US20020032315A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15,
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Best Local (
                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/056,160B
FILING DATE: 06-Apr-1998
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/054,856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1093R2
                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: 491
TYPE: PRT
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                           ZIF: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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APPLICANT:
                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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ADDRESSE: Ge...
1 DNA Way
San Francisco
                                                                                           TYPE: Amino Acid
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                                                                                                            LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24 DIQLTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPGKAPKLLIYFTSSLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS 60
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Similarity 88.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09056160B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Baca, Manuel
Wells, James A.
Presta, Leonard G
Lowman, Henry B.
Chen, Yvonne M.
                                                                                                              107 amino acids
                                                                           Linear
   88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 507; DB 13;
Pred. No. 1.1e-35;
7; Mismatches 6;
   Score 505; DB 9;
Pred. No. 3.6e-36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTIBODIES
                  Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 491;
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                                                                                                                                                                                                   ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-234-671-15
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US-10-234-671-15
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                                                                                                                                Query Match 88.9
Best Local Similarity 88.8
Matches 95; Conservative
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                                                                                                                                                                                                                                                                                 TELEPHONE: 650/225-8674
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/056160

PILING DATE: 06-APR-1998

APPLICATION NUMBER: 60/126446

PILING DATE: 07-APR-1997

APPLICATION NUMBER: 60/054856

PILING DATE: 06-AUG-1997

ATTORNEY/AGENT INFORMATION:

NAME: Cui, Steven X.

REGESTRATION NUMBER: 44,637

REFERENCE/DOCKET NUMBER: P1093R2C1

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
CLASSIFICATION: Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb !
COMPUTER: IHM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Baca, Manuel
 61
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                                                            95;
                      RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVBIK 107
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RFSGSGSGTDYTLTISSLOPEDFATYYCQQYSTVPWTFGQGTKVEIK 107
                                                                                                                                                                                                                                                  LENGTH: 107 amino acids TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94080
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STATE: California
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STREET: 1 DNA Way
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Presta, Leonard G.
Lowman, Henry B.
Chen, Yvonne M.
                                                                                                                                                 88.9%;
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                                                                                                                              Score 505; DB 14; Length 107; Pred. No. 3.6e-36; 6; Mismatches 6; Indels
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RESULT 15
US-10-234-671-103
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US-09-056-160B-105
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                                                                                                                                         Sequence 103, Application US/10
Publication No. US20030190317A:
GENERAL INFORMATION:
APPLICANT: Baca, Manuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 105, Application US/09056160B Patent No. US20020032315A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/054,856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1093R2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb fl.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/056,160
FILING DATE: 06-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: Amino Acids
TOPOLOGY: Lin-
056-160-
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LENGTH: 110 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Presta, Leonard G.
APPLICANT: Lowman, Henry B.
APPLICANT: Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 424
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Baca, Manuel
APPLICANT: Wells, James A.
APPLICANT: Presta, Leonard
APPLICANT: Lowman, Henry B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 88.7%;
Local Similarity 87.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
               NUMBER OF SEQUENCES: 1: CORRESPONDENCE ADDRESS
                                                 Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
                                                                                                                                                                                                                                                                                              61
                                                                                                                                                                                                                                                                                                                   61 RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
                                                                                                                                                                                                                                                                                                                                                                                                                                            94;
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                                                                                                                                                                                                                                                                                                                                                                   DIQLTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPGKAPKVLIYFTSSLHSGVPS
ADDRESSEE: Genentech, Inc
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                                                                                                                                                                                 Application US/10234671 o. US20030190317A1
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                                                                                                         Presta, Leonard
                                                                                                                        Wells, James A.
                                                                                       Lowman, Henry B.
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Pred. No. 4.5e-36;
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STREET: 1 DAN WAY
CITY: SCALL SAN FARCASCO
STATE: CALIFORNIA
COUNTRY: USA
COUNTRY: USA
COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch. 1.44 Mb floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch. 1.64 Mb floppy disk
COMPUTER: 15M PC compatable
OMERATION SYSTEM, PC-DOS/MS-DOS
SOFTWARE MinPatin (Genencech)
APPLICATION MOMBER: US/10/234,671
APPLICATION MOMBER: US/10/234,671
APPLICATION MOMBER: 03-6sp-2002
APPLICATION MOMBER: 03-6sp-2002
APPLICATION MOMBER: 05/056160
APPLICATION MOMBE
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being primand is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2004 Compugen Ltd.
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g kappa chain V-I
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g kappa chain V-I
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g kappa chain V-I
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(C;Species: Homo sapiens (man)
(C;Date: 24-Apr-1984 #sequence_revision 02-Jul-1998 #text_change 09-Jul-2004
(C;Accession: A91653; A01862; S02573
(R;Schiechl, H.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 353, 345-370, 1972
(A;Title: Die Primaerstruktur einer monoklonalen Immunglobulin-L-Kette vom kappa-Typ, A;Reference number: A91653; MUID:72189444; PMID:5028201
A;Accession: A91653
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A;Map position: 2p12-2p12
C;Complex: an immunoglobulin heterotetramer subunit consists of two identical light () hain disulfide bonds; in some cases, such as IgA and IgM, the subunits associate into C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Contents: annotation; X-ray crystallography
A;Note: the structure of the V region was determined by molecular replacement methods u
R;Steiner, V:, Chang, J:Y.
FEBS Lett. 222, 6-10, 1987
A;Title: Chemical modification of the carboxyl groups of protein substrates enhances th
A;Reference number: S02572; MUID:88005152; PMID:3115831
A;Contents: annotation
C;Comment: This is a Bence Jones protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: protein
A;Residues: 1-108 <SCH>
A;Residues: 1-108 <SCH>
A;Residues: 1-108 <SCH>
A;Residues: 1-108 <SCH>
A;Residues: 1-108 <SCH
A;Roser-federences: UNIPROT:P01594
A;Note: the C region of this chain has the Inv (3) marker
R;Pehlhammer, H.; Schiffer, M.; Epp, O.; Colman, P.M.; Lattman, E.E.; Schwager, P.; Ste
Biophys. Struct. Mech. 1, 139-146, 1975
Biophys. Struct. Mech. 1, 139-146, 1975
A;Title: The structure determination of the variable portion of the Bence-Jones protein
A;Reference number: A90729; MUID:77022433; MID:1234024
                                                             밁
                                                                                                                                                                                                                                                                                                     C;Keywords: heterotetramer; immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                  F;23-88/Disulfide bonds: #status predicted
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                                                                                     RFSGGSSTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR
                                                                                                                                                      DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
                                                                                                                                   DIQMTQSPSSLSASVGDRVTITCQASQDISDYLNWYQQKPGKAPKLLIYDASNLESGVPS
                                                           RESGGGSGAHETETISSLOPEDIATYYCOOYDYLPWTFGOGTKVEIKR
                                                                                                                                                                                                           Conservative
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S44122
C26405
K1HURY
S19970
S3627
B48677
B28044
PL0282
S31998
S36264
A28044
A28044
D48677
                                                                                                                                                                                                           5; Mismatches
                                                                                                                                                                                                                            Score 480; DB 1;
Pred. No. 1.7e-36;
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                                                                                                                                                                                                                                             Length 108
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kappa chain V r
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Scoring table: Sequence: Title: Perfect score:

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US-10-089-500-54 568

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Ig kappa chain V region (Py20) - mouse (Species: Mus musculus (house mouse) (;Species: Mus musculus (house mouse) (;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 09-Jul-2004 C;Accession: A38740 R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R. J. Biol. Chem. 266, 6607-6613, 1991 A;Title: Heavy and light chain variable region sequences and antibody proper A;Reference number: A38740; MUID:91177923; PMID:1706720 A;Accession: A38740
Ig kappa chain V region (Py69) - mouse (C,Species: Mus musculus (house mouse) C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_C;Accession: G38740 R,Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R J. Biol. Chem. 266, 6607-6613, 1991 A;Title: Heavy and light chain variable region sequences A;Reference number: A38740; MUID:91177923; PMID:1706720
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G38740
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin
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A;Title: Heavy and light chain variable region se A;Reference number: A38740; MUID:91177923; PMID:1A;Accession: E38740
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C;Superfamily: immunoglobulin V region; immunoglobulin homolC;Keywords: heterotetramer; immunoglobulin
E;19-93/Domain: immunoglobulin homology <IMM>
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A;Molecule type: mRNA
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81.5%;
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11; Mismatches 9
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Pred. No. 2.2e-36;
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A;Experimental source: MLR-lpr/lpr
A;Note: sequence extracted from NCBI backbone
R;Puccetti, A.; Koizumi, T.; Migliorini, P.; P
J. Exp. Med. 171, 1919-1930, 1990

A;Title:

immunoglobulin

light

chain

from

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lupus-prone

mouse ٠ :

induces

autoantibodies

Barrett,

K.J.;

Schwartz

e (NCBIP:60876)
Andre-Schwartz,

A;Status: preliminary; A;Molecule type: nuclei A;Residues: 1-107 <KOI;

y; not compared leic acid

with

conceptual

1-107 <KOI>

A; Accession: B49026

A; Title: Molecular heterogeneity of auto-anti-idiotypic A; Reference number: A49026; MUID:91364791; PMID:1909645

Eur. J. Immunol. 21, 2185-2193, 1991 R; Koizumi, T.; Puccetti, A.; Migliorini,

P.;

Barrett,

K.J.; Schwartz,

R.S.

antibodies

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MLR-lpr/lpr

mice

#text_change

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C;Superfamily: in C;Keywords: hete F;19-93/Domain:
B49026
Ig kappa chain V region, anti-idiotypic monoclonal antibody C;Species: Mus musculus (house mouse)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_cha
C;Accession: B49026; PLO220
C;Accession: B49026; PLO220
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J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences
A;Reference number: A38740; MUID:91177923; PMID:1706720
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C; Date: 18-Oct-1991 #sequence_revision
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A; Residues: 1-111 < RUF>
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Keywords: heterotetramer; immunoglobul;
19-93/Domain: immunoglobulin homology;
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Mus musculus (house mouse)
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Pred. No. 9.4e-36;
.0; Mismatches 10
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Pred. No. 1.8e-35;
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Eur. J. Immunol. 23, 3248-3271, 1993
A,Title: Expressed human immunoglobulin chi
A,Reference number: S40312; MUID:94080891; E
A,Accession: S40367
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KVMS73
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Eur. J. Immunol.
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S40367
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A; Experimental source: (C; Superfamily: immunoglo
                            A; Molecule type: protein A; Residues: 1-108 < SCH>
                                                          A;Title: Determination of the A;Reference number: A01926; MA;Accession: A01926
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Matches 88
Cross-references: UNIPROT:P01643; Comment: This chain was isolated
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;Species: Homo sapiens (man)
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;Accession: PL0220
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Status: nucleic
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Immunol. 23, 3248-3271, 1993
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lobulin V region; immunoglobulin homo
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82.2%; Pred. No. 3.1e-35;
tive 10; Mismatches 9
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Pred. No. 8.6e-35;
6; Mismatches 13
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R;Palm, W.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 356, 167-191,
A;Title: Die Primaerstruktur einer kristalliner
vollstaendige Aminosaeurssequenz des Proteins
                                                                                                             Ig kappa chain V-I region (Rei) - humar C;Species: Homo sapiens (man) C;Date: 24-Apr.1984 #sequence_revision C;Accession: A91663; A01873
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K1HURE
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A; Reference number: A91663; A; Accession: A91663
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A;Residues: 1-108 <WYS>
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78.7%;
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Pred. No. 9e-35;
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                 PMID:809329
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A,Note: the C region of this chain has the Inv (1,2) marker R;Epp, O.; Lattman, E.E.; Schiffer, M.; Huber, R.; Palm, W. Biochemistry 14, 4943-4952, 1975
A,Title: The molecular structure of a dimer composed of the variable portions A;Reference number: A90392; MUID:76039968; PMID:1182131
A;Contents: annotation; X-ray crystallography, 2.0 angstroms C;Comment: This is a Bence Jones protein.
C;Genetics:
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A;Note: the authors translated the codon TC for residue 1 as Leu
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;30-104/Domain: immunoglobulin homology
Ig kappa chain (clone KL2.29 / KL2.33 / KL3.8) - C;Species: Mus musculus (house mouse) C;Date: 14-Feb-1997 #sequence revision 13-Mar-199 C;Accession: S69903; S69904; S69905 R;Wysocki, L.J.; Creadon, G.; Lehmann, K.R.; Camb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ig kappa chain precursor V region (AC-1001) - mouse (fragme: C;Species: Mus muscullus (house mouse)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_ch.
C;Accession: A29380
R;Chen, H.T.; Kabat, E.A.; Lundblad, A.; Ratcliffe, R.M.
J. Biol. Chem. 262, 13579-13583, 1987
A;Title: Nucleotide and translated amino acid sequences of A;Reference number: A92612; MUID:88007582; PMID:3115981
A;Accession: A29380
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A; Residues: 1-108 < PAL>
A; Cross-references: UNIP
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A; Residues: 1-122 < CHE>
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C;Keywords: heterotetramer
F;16-90/Domain: immunoglobu
F;23-88/Disulfide bonds: #s
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
mes 87; Conserv
                                                                                                                                                                                                                                                       61
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                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                  DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DIQMTQSPSSLSASVGDRVTITCQASQDIIXYLNWYQQTPGKAPKLIIYEASNLQAGVPS
                                                                                                                                                                                                                                                                                                     DIQMTQTTSSLSASLGDRVTISCRASQDINNYLNWYQQKPDGTVKLLIHYTSRLHSGVPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                      Score 454; DB 2;
Pred. No. 4.3e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 2.56
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 456; DB 1;
Pred. No. 2.5e-34;
                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
  K.R.; Cambier, J.C.
                                           13-Mar-1997 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mouse (fragment)
                                                                                                mouse
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RESULT 13
JL0080
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A;Status: preliminary; translation not
A;Molecule type: DNA
A;Residues: 1-108 <WYS>
A;Cross-references: EMBL:X55044; NID:g5
                                                                                                                                                                                                                                         C;Keywords: immunoglobulin
F;1-6/Domain: signal sequence (fragment) #status [F;1-6/Domain: signal sequence (fragment) #status predicted
F;7-115/Product: Ig light chain #status predicted
F;22-96/Domain: immunoglobulin homology <IMM>
F;30-40/Region: complementarity-determining 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Accession: S69904
A;Status: preliminary; translation
A;Molecule type: DNA
A;Residues: 1-108 <WY2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunology 75, 116-121, 1992
A;Title: B-cell proliferation initiated by Ia cross-linking and sustained by interleukin A;Reference number: 869900; MUID:92165291; PMID:1537587
                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:M27793; NID:g197161; PIDN:AAA38937.1; PID:g197162 A;Note: the authors translated the codon AGG for residue 30 as Ser C;Superfamily: immunoglobulin V region; immunoglobulin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-108 < WY3 >
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A; Residues: 1-115 < KAA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: EMBL:X55045; NID:g511031;
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67
                                 61 RFSGGGSGTDYTLTISSLOPEDIATYFCHQYSKLPWTFGQGTKVEIKR 108
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                                                                                                         1 DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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                                                                                                                                                                  Similarity
                                                                        DIQMTQTTSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
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                                                                                                                                                 79.6%; Score 452; DB 2;
79.6%; Pred. No. 6.2e-34;
tive 11; Mismatches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        79.6%; Score 452; DB 2; 79.6%; Pred. No. 5.8e-34;
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RESULT 14 S52789 Ig kappa chain V region

human (fragment)

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A;Status preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-129 <ROC>
A;Cross-references: EMBL:X85995; NID:g758588; PIDN:CAA59987.1; PID:g758589
A;Cross-references: EMBL:X85995; NID:g758588; PIDN:CAA59987.1; PID:g758589
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-110 <HAW>
A;Residues: 1-110 <HAW>
A;Residues: 1-100 <HAW>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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R;Rocca, A.; Khamilchi, A.A.; Touchard, G.; Mougenot, B.; Ronco, P.; Denoroy, L.; Deret, submitted to the EMBL Data Library, March 1995
A;Description: Light Chain V region gene usage restriction and peculiarities in myeloma.
Search completed: December 23, 2004, 19:06:05 Job time: 6.83589 secs
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A;Description: Idiotypic vaccination against human B-cell lymphoma: rescue of variable
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S44118
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A; Accession: S52789
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Best Local S
Matches 87
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;Species: Homo sapiens (man)
;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 24-May-2001
;Accession: S44118
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Best Local :
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87; Conserv
                                                                                                                                                                                                                                                                                                                                            h 79.2%; Score 450; DB 2; Length 110; Similarity 80.9%; Pred. No. 8.9e-34; B9; Conservative 6; Mismatches 13; Indels
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                                                                                                           RESGSGSGTDFTFTISSLQPEDIATYYCQQYDNLPPGVTFGPGTKVDIKR 110
                                                                                                                                                     RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLP--WTFGQGTKVEIKR 108
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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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  seq length: 0
seq length: 2000000000
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Match
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Gapop 10.0 ,
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568
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  DIQMTQSPSSLSASVGDRVT.....HQYSKLPWTFGQGTKVEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Q6GMX8
AAR11052
KV1H_HUMAN
Q6GMX0
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KV1K_HUMAN
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KV5U MOUSE
Q7Z3¥4
Q9JL84
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QBRO62
KV10 HUMAN
AAR10992
KVSK MOUSE
KVSK MOUSE
KVSM MOUSE
KV1A HUMAN
KVSO MOUSE
Q9UL77
KVSN MOUSE
Q9UL77
KVSN MOUSE
                                                                                           KV1R_HUMAN
Q91WF8
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KV1Y HUMAN
Q9UL70
                                                    KV1M_HUMAN
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P01593 homo sapien
P01648 mus musculu
P01593 homo sapien
P01648 mus musculu
Q9ul77 homo sapien
P01608 homo sapien
P01608 homo sapien
P01610 homo sapien
Q9ul70 homo sapien
P01610 homo sapien
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REDIJINE=22388257; PubMed=12477932;

RC STRAIN=FVB/N; TISSUE=Colon;

RC MEDIJINE=22388257; PubMed=12477932;

RX MEDIJINE=22388257; PubMed=12477932;

RX MEDIJINE=22388257; PubMed=12477932;

RX MILESCHUL S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,

RA Alteschul S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,

RA Alteschul S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,

RA HOpkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Karywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Mones S.J., Marra M.A.,

"Generation and initial analysis of more than 15,000 full-length human

RT and mouse cDNA sequences.",

Proc. Matl Anada Sci II S. A Gallaga-1600/2002
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Mammalia; Eutheria;
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Mus musculus (Mouse).
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01-MAR-2004
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                                             Acad. Sci.
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Pred. No. 9
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                                             99:16899-16903 (2002)
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KV5J MOUSE STANDARD; PRT
P01643;
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05-JUL-2004 (Rel. 44, Last annota
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EMBL; BC013496; AAH13496.1; -.
PIR; A38740; A38740.
PIR; C38740; E38740.
PIR; B38740; G38740.
PIR; G38740; G38740.
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                                                                                                                                                                                                                                                                                                                                                                                  "Determination of the primary structure of immunoglobulin. Amino-acid sequence of the Eur. J. Biochem. 59:525-337(1975).

-I- MISCELLANEOUS: This chain was isolated PIR; A01926; KVMS73.

HSSP; P01594; 1JV5.
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PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
                                                                                                                                                                                           DOMAIN
DOMAIN
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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rrPro; IPR003597; Ig_cl.
rrPro; IPR003597; Ig_MC.
rrPro; IPR003596; Ig_v.
rrPro; IPR003596; Ig_v.
rrPro; IPR003596; Ig_v.
rrpro7654; Cl-set; l.
rrpr00047; Ig; l.
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Framework-4.
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RC STRAIN=FVB/N, TISSUB=Colon;
RX MEDLINE=22388257; PubMed=12477932;
RX MEDLINE=22388257; PubMed=12477932;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettenan M., Madan A., Rodrigues S., Sanchez A.,
RA Milting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mitting M., Madan A., Young A.C., Shevchenko M.C.,
RA Rodriguez A.C., Grimwood J. W., Green E.D., Dickson M.C.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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                                                                                                                                                                                              InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF007654; Cl-Bet; I.
Pfam; PF00767; Ig; 1.
SMART; SM00406; IGy; 1.
SMART; SM00406; IGy; 1.
PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS50835; IG_MHC; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                              Strausberg R.;
Submitted (APR-2002) to the
EMBL; BC027418; AAH27418.1;
HSSP; P01594; 1JV5.
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Mus musculus (Mouse).
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01-JUN-2002 (TrEMBLrel.
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STRAIN=FVB/N; TISSUE=Colon;
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                             Hypothetical
SEQUENCE 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Generation and initial analysis of more than 15,000 and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
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                                                                                                                                                      al protein.
234 AA; 25858 MW;
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illarity 79.6%;
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                                                                    Score 461; DB
Pred. No. 2.4e
.3; Mismatches
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3; Mismatches
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-I- MISCELLANEOUS: This is a Bence-Jones PIR; A91663; K1HURE.
PDB; 1AR2; X-ray; @=1-107.
PDB; 1RW; X-ray; @=1-107.
PDB; 1REI; X-ray; A/B=1-107.
PDB; 1REI; X-ray; A/B=1-107.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0003823; F:immune response; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR003506; Ig_v.
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DOMAIN 89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kappa-type I-chain, subgroup I (Bence-Jones protein Rei); isolatic and characterization of the tryptic peptides; the complete amino sequence of the protein; a contribution to the elucidation of the three-dimensional structure of antibodies, in particular their combining site.";
                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00047; ig; 1.
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Palm W., Hilschmann
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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DT 05-JUL-1986 (Rel. 01, Last sequence update)
DT 05-JUL-2004 (Rel. 44, Last annotation update)
DE 1g kappa chain V-V region HP R16.7.
S Mus musculus (Mouse).
C Eukaryota, Metazoa; Chordata; Craniata; Vertebrat C Mammalia; Eutheria; Rodentia; Sciurognathi. Mouse.
NCBI TaxID=10090;
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02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musimus.
                                                                                                                                                                                                               MOUSE
KV5K_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liang Z., Xie C., Chen C., Kreska D., Hsu K., "Antinuclear autoantibodies from B6.Slel mice. Submitted (SEP-2003) to the EMBL/GenBank/DDBJ EMBL; AY436832; AAR10992.1; -.
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STRAIN=B6.Sle1; TISSUE=Spleen;
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Pred. No. 5
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Pred. No. 3.2e-40;
7; Mismatches 14
                                            Craniata; Vertebrata; I
Sciurognathi; Muridae;
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                                                                   Euteleostomi;
                                              Murinae;
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RESULT 8

KV5L WOUSE

ID KV5L WOUSE

AC P01645;

P01645;

P01645;

P01645;

P017 21-JUL-1986;

P07 21-JUL-1986;

P07 21-JUL-1986;

P08 20-JUL-2004;

P08 Mus musculus

OC Mammalla; But

OX NCBI_TAXID=10

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RN [1]

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RC STRAIN=A/J;

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RA Siegelman M.,

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DR HSSP; P01607;

DR InterPro; IPF

DR Ffam; P80044

SMART; SMO44

DR SMART; SMO44

DR SMART; SMO44

DR PROSITE; P856;

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HSSP; P01594; 1JV5.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
                                                                             -i- MISCELLANEOUS: Anti-arsonate
HSSP; P01607; 1REI.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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STRAIN=A/J;
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44, Last annotation updat
V region HP 93G7.
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Rodentia;
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79.6%;
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Pred. No. 3
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Complementarity-determining-3.
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By similarit
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                   Complementarity-determining-1.
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Sciurognathi;
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2.3e-39;
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thi; Muridae; Murinae; Mus
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Best Local :
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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SMART; SM00406; IGv;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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PS50835; IG_LIKE;
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RFSGSGSGTDYSLTISNLEQEDIATYFCQQGYMLPRTFGGGTKLEIKR
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01, Last sequence up
44, Last annotation
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80.6%;
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79.6%;
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Pred. No. 2.9e-
8; Mismatches
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Pred. No. 2.9e-39;
0; Mismatches 12
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Framework-3.
Complementarity-determining-3.
Framework-4.
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                                                                     SNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
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2.9e-39;
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Best Local S
Matches 85
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HSSP; P01607; 1BWA:
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0005823; F:antigen binding; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
Pfam; PF00047; 19; 1.
SMART; SM00406; IGV; 1.
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21-JUL-1986 (Re
21-JUL-1986 (Re
05-JUL-2004 (Re
1g kappa chain
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21-JUL-1986
21-JUL-1986
05-JUL-2004
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                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                Mus musculus (Mouse)
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KV50_
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DOMAIN
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-i- MISCELLANEOUS: The C region of this chain has the INV-I- MISCELLANEOUS: This is a Bence-Jones protein.
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Mammalia; Eutheria; Primates;
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ein; Direct
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44, Last annotation updat
V region HP 91A3.
                                                                              Chordata;
Rodentia;
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Pred. No. 4.7e-39;
9; Mismatches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Complementarity-determining-3 Framework-4.
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Complementarity-determining-2
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                                                                          Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Catarrhini; Hominidae;
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SEQUENCE. STRAIN=A/J;

NCBI_TaxID=10090; [1]

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RESULT 12
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Best Local S
Matches 84
Query Match
                                                                                         PIR; B49047; B49047.

PIR; S34083; S34083.

HSSP; PO1607; IBWW.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

Pfam; PF00047; ig; 1

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.
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DOMAIN
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MEDLINE=98277139; PubMed=9614934;
WW X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Complete amino acid sequence of light chain variable regions derived from five monoclonal anti-p-azophenylarsonate antibodies differing with respect to a crossreactive didotype.";

Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-OCT-2003
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01-MAY-2000
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PROSITE; PS50835; IG_LIKE;
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SMART; SM00406; IGv;
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HSSP; P01594; 1475.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                 EMBL; AF035037; AAD56273.1;
                                                                                                                                                                                                                                                                                                                                              Young
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Myosin-reactive
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                                                                                                                                                                                                                                                                                                                           "Myosin-reactive autoantibodies
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                                                                                                                                                                                                                                                                                   Immunol. Immunopathol.
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Ig_v.
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Last annotation update)
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Pred. No. 4.7e-39;
                                                                                                                                                                                                                                                                                   87:184-192(1998).
  Score 445;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                         C06681716C4D16F3
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KV5N_MOUSE
ID KV5N
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Best Local S
Matches 85
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Matches 86; Conserv
              Q6GMW1;
Q6GMW1;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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P01647;
21-JUL-1986
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NON TER
SEQUENCE
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InterPro; IPR0071110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUL-1986 (Rel. 01, Creat
21-JUL-1986 (Rel. 01, Last
05-JUL-2004 (Rel. 44, Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               with respect to a Proc. Natl. Acad
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=A/J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plete amino acid sequence of light chain variable regions five monoclonal anti-p-azophenylarsonate antibodies differespect to a crossreactive idiotype.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61
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                                                                                                                                                                                                RFSGSGSGTDYSLTISNLEQEDIATYFCQQGKTLPRTFGGGTKLEIKR
                                                                                                                                                                                                                                     RFSGGGSGTDYTLTISSLQPEDIATYFCHQYSKLPWTFGQGTKVEIKR
                                                                                                                                                                                                                                                                              DIOMIQTISSISASIGDRVTISCRASQDINNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
                                                                                                                                                                                                                                                                                                    DIQMTQSPSSLSASVGDRVTITCSASQDISNYLNWYQQKPDKAVKLLIFYSSNLHSGVPS
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(TrEMBLrel.
(TrEMBLrel.
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Acad. Sci. U.S.A. 78:7679-7683(1981).
protein
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35 56
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89 108
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el. 44, Last annotation
V-V region HP 124E1.
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78.7%;
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Pred. No. 7.7e
10; Mismatches
                  Last sequence update)
Last annotation update)
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                                                                                                  PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              similarity.
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.7e-39;
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RESULT 15

KV1P HUMAN

ID KV1P HUMAN

STANDARD;

AC P01608;

AC P01608;

AC P01508;

AC P01508;

AC P01608;

AC P01608;

AC P01608;

C 21-JUL-1986 (Rel. 01, Create DT 21-JUL-1986 (Rel. 04, Last e DT 05-JUL-2004 (Rel. 44, L
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DR InterPro; IPR007110; Ig-1ike.

DR InterPro; IPR003597; Ig-C1.

DR InterPro; IPR003596; Ig-MHC.

DR InterPro; IPR003596; Ig-MHC.

DR Ffam; PP00647; Ig: 2.

DR SMART; SM00409; IG; 2.

DR SMART; SM00407; IGC1; 1.

R SMART; SM00406; IGV; 1.

R SMART; SM00406; IGV; 1.

R PROSITE; PS50835; IG LIKE; 2.

PROSITE; PS50835; IG LIKE; 2.

PROSITE; PS50835; IG LIKE; 2.

PROSITE; PS50835; IG-MHC; UNKNOWN 1.

Hypothetical protein.

SEQUENCE 236 AA: 75777
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buctow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh P.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh P.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh P.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

A Hopkins R.F., Jordan H., Farmer A.A., Rubin G.M., Hong L.,

A Brownstein M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Brownstein M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Caraninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Hichards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Hahting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

A Jones S.J. May A.
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Best Local S
Matches 87
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Eukaryota; Metazoa; (
Mammalia; Eutheria; I
NCBI TaxID=9606;
[1]
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Submitted (JUN-2004) to the
EMBL; BC073791; AAH73791.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. TISSUE-Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jones S.J., Marra M.A.; "Generation and initial analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Spleen;
MEDLINE-22388257; PubMed-12477932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                  (Rel. 01, Created)
(Rel. 01, Last sequence up
(Rel. 44, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
    Chordata;
Primates;
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    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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GO; GO:0005823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-1ike.
InterPro; IPR003596; Ig-v.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Bence-Jones protein; Direct protein sec
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PIR; A91638; K1HURY.
HSSP; P01607; 1BWW.
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Complementarity-determining-3.
Framework-4.
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Complementarity-determining-2.
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SUMMARIES

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Ganglioside 03-JUL-2001 AAB81989; AAB81989 standard; protein; GD3 (first entry) specific antibody related protein SEQ ID NO: 119 AA

ALIGNMENTS

Ganglioside; GD3; complementarity determining region; CDR; antibody; cancer.

Mus musculus.

WO200123432-A1.

05-APR-2001.

29-SEP-2000; 2000WO-JP006774

30-SEP-1999; 99JP-00278291. 06-APR-2000; 2000JP-00105088.

(KYOW) KYOWA HAKKO KOGYO KK.

Hanai N, Shitara K, Nakamura ζ, Niwa

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WPI; 2001-266143/27.

New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.

Claim 10; Page 173-174; 183pp; Japanese.

The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention

Sequence 119 AA;

Query Match 100.0%; Score 624; DB 4; I Best Local Similarity 100.0%; Pred. No. 7.4e-51; Matches 119; Conservative 0; Mismatches 0; Length 119; Indels 0 Gaps 0

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RESULT 2
ABULIOL2
ID ABULL
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XX ABULL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drugs containing genetically-modified antibody against ganglioside GD3, its fragment, immunocompetent cell activators or/and antitumor agents in combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance acapable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 7; Page 112-113; 121pp; Japanese.
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                                                                                                                                                                                                                                                                                                                              EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SDSVKGRFTISRDNAKWTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
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                                                                                                                                                                                      SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                    protein; 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                            Score 624; DB 6;
Pred. No. 7.4e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ۲,
                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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RESULT 4
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Best Local
          Monoclonal antibody; Ab; ganglioside GM2; chimera; chimeric antibody; expression vector; heavy; light; chain; hypervariable region; CDR; constant region; hybridoma; Ig; immunoglobulin; promoter; enhancer.
                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAR33256-57 represent rat heavy and light chain variable regions respectively. The DNA sequences encoding these proteins were used in the construction of humanised chimeric antibody expression vectors. In these humanised antibodies none of the amino acids of the non-human animal Ab variable region have been changed. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Promoter; v
humanised;
                                                                                                                                                                                                                                                                                                                                       Sequence 130 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Humanised chimeric antibody prodn. against ganglioside GD3 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAQ33257.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rattus rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rat immunoglobulin H chain varible region of pKM641HA3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
12-JUL-1993
                                                          KM641 L
                                                                                  18-NOV-1994
                                                                                                                                AAR53341 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1993-095510/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAR-1993.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein
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                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                   SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                        EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 29-30; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        such as melanoma,
                                                                                                                                                                                           SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      variable; region; rat; immunoglobulin; heavy;
; chimeric; antibody; expression vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanai N,
                                                                                                                                                                                                                                                                                          Conservative
                                                                                 (first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HAKKO
                                                          variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91JP-00238375
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11. .130
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                                                                                                                               protein; 130
                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hasegawa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroblastoma,
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                                                                                                                                                                                                                                                                                        Score 624; DB 2;
Pred. No. 8.2e-51;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide"
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Mus musculus

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RESULT 5
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XX AAY2
XX AAY2
XX PON
DT 04-N
DT 04-N
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         monoclonal Ab KM641-producing cells was isolated and KM641 H and L chain cDNAs isolated. The base sequences of the Ig variable regions in KM641 H chain cDNA (pKM641HA3) and KM641 L chain cDNA (pKM641HA3) and KM641 L chain cDNA (pKM641HA2) are given in AAQ45438-9. A KM641-derived chimeric human Ab H chain expression vector pKM641HA3 to a vector for chimeric human bh H chain expression using the synthetic DNAs given in AAQ63439 and AAQ63440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Humanised cytocidal
  07-SEP-1993;
                                                                                                                                                                                                                                                                                                                   pKM641 HA3 immunoglobulin heavy chain
                                                                                                                                                                                                                                                                                                                                                                   04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                      AAY28369;
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                                                07-JUN-1995;
                                                                                                17-AUG-1999
                                                                                                                                              US5939532-A
                                                                                                                                                                                           Mവ8 8p.
                                                                                                                                                                                                                                              light
                                                                                                                                                                                                                                                                     antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-SEP-1992;
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                                                                                                                                                                                                                                            ody; nucleotide; genomic; hypervariable region; chimeric;
chain; amino acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody specific for ganglioside GM2 - used for producing a effect on cancers such as melanoma, neuroblastoma and glioma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 116-117; 191pp; English
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Pred. No. 8.2e-51;
Mismatches 0;
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RESULT 6
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Best Local S
Matches 119
WPI; 2000-402204/35.
N-PSDB; AAA51003.
                                                                               18-SEP-1991;
18-SEP-1992;
                                                                                                                                                                                              Protein
                                                                                                                                                                                                                                 Key
                                                                                                                                                                                                                                                                                                               Murine immunoglobulin heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             no methionine initiation codon and the leader sequence was partly lacking. The chimeric human antibodies are useful in the treatment of cancer, especially that which is of mouse meral ectodermal origin. In contrast to prior art constructs based on mouse menoclonal antibodies, the chimeric human antibodies do not cause anti-mouse immunoglobulin production. The chimeric human antibodies have a prolonged half-life and
                                                                                                                                       28-JUN-2000
                                                                                                                                                                                                                                                       Mus sp.
                                                                                                                                                                                                                                                                                                                                                             AAB01627;
                                                        (KYOW)
                                                                                                                                                             EP1013761-A2
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a reduced frequency
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                                                                                                                                                                                                                                                                                        immunoglobulin; H chain; heavy chain; variable region; cancer;
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                                                        куома накко кодуо
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                                  Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                              91JP-00238375.
92EP-00116026.
                                                                                                                99EP-00124345
                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                /label= mature_immunoglobulin_heavy_chain_variable region
                                                                                                                                                                                                         'label= signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of adverse effects when compared to mouse monoclonal
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                                 Hasegawa
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Pred. No. 8.2e-51;
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                                Miyaji H,
                                  Kuwana
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Matches 119
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   The present invention describes a monoclonal antibody which can react specifically with gaughioside GD3. The antibody and its derivatives a useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplificat of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antibodies as they do not provoke a reaction in the human and side effects, such as the formation of anti-mouse immunoglobulin antibody the rapid half-life of the immunoglobulins, do not occur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is a murine immunoglobulin heavy chain variable region from plasmid KW-641. The coding sequence was used in the creation of an expression vector, along with the sequence for a human antibody, to produce humanised chimaeric antibodies, which can be used to treat cancer. Humanised chimaeric antibodies are more effective than mouse
                                                                                                                                                                                                                                                               New human type complementation-determining region-transplanted and derivatives against ganglioside GD3, useful in diagnosis ar of e.g. tumors, with low antigenicity, little side effects but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 14; Page 27-28; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ganglioside;
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                                                                                                                                                                             Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Shitara K,
                                                                                                                                                                             Page 138-139;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GD3
                                                                                                                                                                                                                                         cancer.
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                                                                                                                                                                          183pp; Japanese.
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Pred. No. 8.2e-51;
; Mismatches 0;
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Best Local S
Matches 119
                                                                                                                                                                                                               Query Match
Best Local S
Matches 119
                                                                                                                                                                                                                                                                                                                                                                 The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumo like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This seque represents a protein associated with the anti- ganglioside GD3 antibod
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Drugs containing genetically-modified antibody against ganglioside its fragment, immunocompetent cell activators or/and antitumor ager combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                        Sequence 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 97; 121pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-067410/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2001; 2001JP-00097483
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU11002 standard;
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                              SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                      EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                       EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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  SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
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Pred. No. 8.7e-51; Mismatches 0
                                                                                                                                                                                                               Score 624; DB 6;
Pred. No. 8.7e-51;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GD3 antibody; tumour; melanoma.
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RESULT 9
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RESULT 10
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Best Local S
Matches 119
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06-APR-2000; 2000JP-00105088
Synthetic
                                             Ganglioside;
                                                                       Ganglioside
                                                                                                        03-JUL-2001
                                                                                                                                                                AAB81985
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119; Conserv
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                                                                                                                                                                standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                          582 AA;
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                                                                      specific antibody related protein SEQ ID NO:
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                                                                                                                                                                protein; 119
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                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 624; DB 4;
100.0%; Pred. No. 4.2e-50;
tive 0; Mismatches 0;
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                                          determining region;
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RESULT 11
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Best Local Sim
Matches 109;
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                                                                                                                                                                                                                                                   29-MAR-2002; 2002WO-JP003170
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06-APR-2000; 2000JP-00105088.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                             Niwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GD3; anti-ganglioside GD3
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Pred. No. 1.2e-46;
5; Mismatches 5
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Drugs containing genetically-modified antibody against ganglioside

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RESULT 12
AAB81987
ID AAB81
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specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
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                                                                                                                                                                                        New human type complementation-determining region-transplanted and derivatives against ganglioside GD3, useful in diagnosis ar of e.g. tumors, with low antigenicity, little side effects but
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                                                                                                                              Claim 41; Page 168-172;
                                                                                                                                                                                                                                                                                      WPI; 2001-266143/27.
                                                                                                                                                                                                                                                                                                                              Hanai
                                                                                                                                                                                                                                                                                                                                                                                                              30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
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                                                                                                                                                                                                                                                                                                                            Nakamura
                                                                                                                              183pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 579; DB Pred. No. 1.2e 5; Mismatches
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.2e-46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     region;
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Sequence 582 AA;

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RESULT 13
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Best Local S
Matches 109
                                                                                       Query Match
Best Local (
                                                                           Matches
                                                                                                                                                  This is the amino acid sequence of the heavy chain variable region of the monoclonal antibody (MAb) for TIA which recognises the human leukocyte differentiation antigen CD6. The MAb is a murine IgG2-type Ab produced by the usual hybridoma techniques. The coding sequence for the variable region can be used to generate humanised variants, especially by mutagenic PCR. CD6 has been shown to be expressed in Tlymphocytes involved in the inflammatory infiltration of the skin in psoriasis. The anti-CD6 MAb can therefore be used to diagnose and treat psoriasis. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                            Sequence 119
                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Heavy chain
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12-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW29994;
                                                                                                                                                                                                                                                                                                                     Monoclonal antibody against human CD6 antigen - useful for diagnosis
                                                                                                                                                                                                                                                                                                                                                                        Sierra Blazquez P,
                                                                                                                                                                                                                                                                                                                                                                                    Montero Casimiro
                                                                                                                                                                                                                                                                                                                                                                                                                                       17-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9719111-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW29994 standard;
                                                                                                                                                                                                                                                                                                                                                                                                              (IMMU-) CENT IMMUNOLOGIA MOLECULAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          eukocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
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                                                                                                                                                                                                                                                                                3; Page 27; 38pp; Spanish
61
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 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                          EVQLVESGGGLVKPGGSLKLSCAASGFKFSRYAMSWVRQTPEKRLEWVATISSGGSYIYY
                                                EVILVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          differentiation antigen;
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             variable region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                        psoriasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variable region; monoclonal antibody; human; CD6; murine; fferentiation antigen; hybridoma; humanisation; mutagenesis; amplification; inflammatory infiltration; skin; psoriasis;
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Pred. No. 7e-46;
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Pred. No. 1.5e-38;
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61

Matches

Best Local Similarity

96;

Conservative

9

Pred. No. 6e-38; 6; Mismatches

17;

Indels

<u>ب</u>

Gaps

60 60

79.3%;

RESULT 15 AAB20442

AAB20442;

21-JUN-2001

(first entry)

AAB20442 standard; protein;

294

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Factor IX; FIX; Factor IXa; FIXa; scFv; antibody; procoagulant; Factor VIII cofactor; blood coagulation disorder; haemophilia A; haemorrhagic diathesis; haemostatic; amidolytic; therapy; mouse;

myc-tag

Anti-FIX/FIXa antibody 198/B1-myc-tag fusion.

밁 S 밁 Ś 片 S

S 119

61 61

PDSVKGRFTISRDNAKNTLYLOMSSLKSEDTAMYYCARLDGYNYRWYFDVWGTGTTVTVS SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTY--YFDSWGQGTTLTVS 118 EVQLVESGGDLVKPGGSLKLSCAASGFTFSRYGMSWVRQTPDKRLEWVATISSGGSYTYY EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY

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Query Match
                           Sequence 134 AA;
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78.0%;

Score 487;

띮 6

Length 134;

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RESULT 14
ABP60555
                                                     The invention relates to a novel monoclonal antibody having the identifying characteristics of, or that is a monoclonal antibody 14F3. An antibody of the invention has osteopathic, antirheumatic, antiarthritic, antiinflammatory, cytostatic, antipsoriatic, antidiabetic, and neuroprotective activity. The polynucleotides encoding the antibodies of the invention may have a use in gene therapy. The antibodies and polypeptides are useful for treating or preventing osteopathic diseases, such as rheumatoid arthritis, osteoporosis, metastatic and primary bone cancer, wear debris induced osteolysis or osteoarthritis, and immune diseases or multiple sclerosis. The present sequence represents the heavy chain variable region of the murine monoclonal antibody 14F3 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse; monoclonal antibody; 14F3; osteopathic; antirheumatic; antiarthritic; antinflammatory; cytostatic; antipsoriatic; antidiabetic; neuroprotective; gene therapy; rheumatoid arthritis; osteoporosis; bone cancer; osteolysis; osteoarthritis; immune disease; psoriasis; insulin-dependent diabetes; inflammatory bowel disease; multiple sclerosis; heavy chain variable region;
                                                                                                                                                                                                                                                                                                 New monoclonal antibody having the characteristics of a monoclonal antibody 14F3, useful for treating or preventing osteopathic diseases, e.g. rheumatoid arthritis, or osteoporosis, and immune diseases e.g.
                                                                                                                                                                                                                                                         Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-MAY-2002; 2002WO-US014246
                                                                                                                                                                                                                                                                                                                                                                                                                                           (SMIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-MAY-2001; 2001US-0292031P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine antibody 14F3 heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAR-2003
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SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                 ABV99887.
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                                                                                                                                                                                                                                                                                       or diabetes
                                                                                                                                                                                                                                                                                                                                                                                                           Tornetta MA,
                                          The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            determining region; CDR.
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50. .66
                                                                                                                                                                                                                                                      8; 51pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= CDR2
99. .110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'label=
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                                             contains
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                                                                                                                                                                                                                                                        English
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                                           three
                                          complementary determining regions
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WPI; 2001-290358/30.
N-PSDB; AAF30732.
                                                                                                                                                                                      Key
Peptide
               Scheiflinger F,
                                                                                          Peptide
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                                                                                                                                                       Region
                                   14-SEP-1999;
                                              13-SEP-2000; 2000WO-EP008936
                                                                 WO200119992-A2
                                                                                                    Protein
                                                                                                               Peptide
                                                                                                                                   Peptide
                                                                                                                                              Misc-difference
                                                                                                                                                                  Protein
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                                                                                                                                                                                                      Chimeric
                                                                                                                                                                                                           Escherichia coli.
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                         BAXTER
                                   99AT-00001576
               Kerschbaumer
                                                                          /label= His_tag
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                                                                                                                                                                                           Location/Qualifiers
                                                                                               label= Myc_tag
                                                                                                                              label=
                                                                                                                                                   label= VH
                                                                                                                                                            label= scFv
                                                                                                                                                                                label= Signal_peptide
                                                                                                                                        note= "encoded
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                                                                                                         label= Spacer
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                                                                                     Spacer
                                                                                                                               Linker
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               Falkner F,
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               Dorner F;
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Search completed: December 23, 2004, 18:57:39 Job time: 33.8339 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of a fusion protein comprising: a PelB CC leader; a single chain Fv (gcFv) derivative of antibody 198/Bl comprising CC the heavy (VH) and 11ght (VL) chain variable regions of 198/Bl joined by CC an artificial, flexible linker peptide; a spacer; a Myc-tag peptide; a cc spacer; and a C-terminal GHis affinity tail. 198/Bl is an example of anti-cc chuman Factor IX (FIX)/activated Factor IX (FIX) antibodies of the CC invention. Anti-FIX/FIX antibodies and their derivatives, including scFv CC fragments, have FVIIIa cofactor activity or FIXa activating activity. CC Administration leads to an increase in the procoagulant activity of FIXa, CC even in the presence of FVIII inhibitors. This allows for rapid blood CC coagulation even in the absence of FVIII or FVIIIa, and in the case of CC laimed pharmaceutical composition for treating patients with blood CC coagulation disorders, especially haemophila A and haemorrhagic CC diathesis. The scrivity especially haemophila A and haemorrhagic CC diathesis. The scrivity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New factor IX/factor IXa antibodies and their derivatives useful for increasing amidolytic activity of factor IXa, and for treating blood coagulation disorders such as hemophilia A and hemorrhagic diathesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 294 AA;
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FVIII-like activity
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                                                                                                                                          143
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                                                                                                                                                                                                                                                                       83 PDSVKGRFTISKDNAKNTLYLQMSSLRSEDTAMYYCTREGGGFTVNWYFDVWGAGTSVTV 142
                                                                                                                                                                                                                                                                                                                  61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLG---TYYYEDSWGQGTTLTV
                                                                                                                                                                                                                                                                                                                                                                                                           ch 78.0%; 1 Similarity 78.7%; 96; Conservative
                                                                                                                                          SS 144
                                                                                                                                                                                                          SS 119
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
   Score
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     Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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624
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     BG
              US-09-225-322B-18
US-09-764-304-18
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US-09-764-304-8
US-08-875-674A-1
PCT-US94-07659-2
US-08-875-674A-3
US-08-879-378A-16
US-08-879-378A-16
US-08-16-335-2
US-09-416-355-2
US-09-416-317-2
US-09-418-737-2
US-09-48-7312-20
US-09-48-7312-20
US-08-497-312-20
US-08-498-3199-16
US-08-488-3199-16
US-08-488-3199-16
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US-08-488-3199-16
US-08-339-582-2
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Sequence 18, Appl Sequence 8, Appli Sequence 8, Appli Sequence 2, Appli Sequence 16, Appli Sequence 16, Appli Sequence 16, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 11, Appli Sequence 11, Appli Sequence 20, Appli Sequence 4, Appli Sequence 16, Appli Sequence 17, Appli Sequence 18, Appli Sequence 18, Appli Sequence 18, Appli Sequence 3, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli
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00 AG W ⁸ 60	RESULT US-09-1	
Query M Best Lo Matches		0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
atch cal s 119 1 11 61	SULT 1 -09-25-32B-18 Sequence 18, Application US/ Sequence 18, Application US/ Sequence 18, Application US/ Sequence 18, Application US/ GENERAL INFORMATION: APPLICANT: SHITARA, KENYA APPLICANT: HANGEANA, MAMOR APPLICANT: KUWANI, NOSUO APPLICANT: KUWANI, NOSUO APPLICANT: KUWANI, YOSHIHI ITILE REFERENCE: 249-101 CURRENT FILING DATE: 1995-05- FRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 1995-05- FRIOR FILING DATE: 1995-05- FRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 1994-08- PRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 1992-09- PRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 1991-09- NUMBER OF SEQ ID NOS: 19 SOFTWARE: PATORE IPRIOR APPLICATION SEG ID NOS 18 LENGTH: 130 TYPE: PRTORANION: Descrip OTHER INFORMATION: Descrip	4 4 4 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6
Similarity 9; Conservat EVTLVESGGDFV EVTLVESGGDFV SDSVKGRFTISR	a derdededed. Pa t	71 5 6 6 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
100. 100. 7ative	Cation US/0922 N: N. KENYA A, KENYA ANA, MAMORU I, HIROMASA A, YOSHIHISA N: PROCESS FOR 49-101 NUMBER: US 08 1995-05-31 NUMBER: US 08 1995-03-21 NUMBER: US 08 1995-03-21 NUMBER: US 08 1995-03-1 NUMBER: US 08 1995-09-17 NUMBER: US 07 1991-09-18 VOS: 19	1443 1119 1129 1129 1136 1136 1245 1247 1247
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Score 624; DB Pred. No. 8.8e 0; Mismatches VSCAASGFAFSHYAMS VSCAASGFAFSHYAMS VSCAASGFAFSHYAMS VSCAASGFAFSHYAMS VLQMRSLRSEDSAMYF YLQMRSLRSEDSAMYF YLQMRSLRSEDSAMYF	5322B PRODUCING HUWA 09/225,322B 15 14404,680 1408,133 1/292,178 947,674 238375	PCT-US96-13152-4 US-08-331-39BA-46 US-08-331-39BA-46 US-08-759-804A-46 US-08-759-804A-46 US-09-227-693-46 US-09-318-786-31 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-33 US-08-976-183A-34 US-09-96-086-5 US-09-96-087-5 5455030-9 US-08-428-197-1 ALIGNMENTS
1.4; Length 130; 1-60; Indels 0; Gaps 0;	NIZED CHIMERA ANTIBODY	Sequence 4, Appli Sequence 46, Appl Sequence 46, Appl Sequence 46, Appl Sequence 31, Appl Sequence 31, Appli Sequence 31, Appli Sequence 31, Appli Sequence 32, Appli Sequence 32, Appli Sequence 32, Appli Sequence 34, Appli Sequence 34, Appli Sequence 5, Appli Sequence 1, Appli Sequence 1, Appli

RESULT 2 US-09-764-304-18 ; Sequence 18, Application US/09764304

Patent No. 6495666

INFORMATION:

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APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT APPLICATION NUMBER: 09/225,322
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1995-03-21
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EARLIER FILING DATE: 1995-03-21
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Best Local S
Matches 119
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/09225322B
Patent No. 6437098
SOFTWARE: PatentIn Ver. EQ ID NO 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1992-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EARLIER APPLICATION NUMBER: JP 3-238375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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WS-08-875-674A-1 US-08-875-674A-1 ; Sequence 1, Application US/08875674A ; Patent No. 6572857

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APPLICANT: HANAI, NOBUS
APPLICANT: HANAI, NOBUS
APPLICANT: MIYAJI, HIROMASA
APPLICANT: MIYAJI, HIROMASA
APPLICANT: MIYAJI, HIROMASA
APPLICANT: MUYANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
PILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-09
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1994-008-17
EARLIER FILING DATE: 1994-008-17
EARLIER APPLICATION NUMBER: US 08/292,178
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, FEATURE:
, OTHER INFORMATION: Description of Artificial Sequence: cDNA KM-641
US-09-225-322B-8
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US-09-764-304-8
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LENGTH: 130
TYPE: PRT
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                                                                                                                                                            Best Local Similarity 99.2 Matches 118; Conservative
                                                                                                                                                                                                Query Match
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EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
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                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                               OTHER INFORMATION: cDNA KM-641
                                                                                                                                                                                                                                                                                  FEATURE:
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71
                   61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
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                                                                          EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAGISSGGSGTYY
SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
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Pred. No. 1.1e-58;
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Pred. No. 1.1e-58;
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RESULT 6
PCT-US94-07659-2
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Patent No. 6572857
OTHER INFORMATION:
OTHER INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 Amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: Protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: -N Terminal fragment.
ORIGINAL SOURCE:
ORGANISM: Mice Balb/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
REGISTRATION NUMBER: 20,910
REFERENCE/DOCKET NUMBER: P-12
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,674A
FILING DATE: 17-Uuly-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Unc
STREET: New York
TATE: New York
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MEDIUM TYPE: Floppy disk 3.5'' (1.4 MB).
COMPUTER: Compatible PC IBM (80486, 8 M Ram).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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OPERATING SYSTEM:
SOFTWARE: Word Pe
                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       IDENTIFICATION METHOD: Experimental.
OTHER INFORMATION: Sequence corresponding to the variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Mice Balb/C INDIVIDUAL ISOLATE: ic TISSUE TYPE: Murine h:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Amino acid
STRANDEDNESS: Un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/CU96/00004 FILING DATE: 18-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
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                                                                                                                                                                                                                              97;
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                                                                                                              SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                            EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60
                                                                          PDSVKGRFTISRDNVKNTLYLQMSSLRSEDTAMYYCARRDYDLDYFDSWGQGTTLTVSS 119
                                                                                                                                                   EVQLVESGGGLVKPGGSLKLSCAASGFKFSRYAMSWVRQTPEKRLEWVATISSGGSYIYY 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  119 Amino acid residues.
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P REZ RODR GUEZ, R.
P REZ RODR GUEZ, P.
                                                                                                                                                                                                                              Conservative
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Word Perfect 5.0 for Windows 95.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TORMO BRAVO, B. R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MONTERO CASIMIRO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lackenbach Siegel Marzullo Aronson & Greenspan, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (914) 723-4300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OLATE: ior tlA
Murine hibridoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unknown.
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                                                                                                                                                                                                                                         79.0%;
81.5%;
                                                                                                                                                                                                                                                                                                                    of the heavy chain of the monoclonal antibody recognizing hudesignated as sub-clone for tlA.
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Pred. No. 1e-45;
3; Mismatches
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                                                                                                                                                              Patent No.
                                                                                                                                                                Sequence 3, Application US/08875674A Patent No. 6572857
                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 77.9%;
Best Local Similarity 79.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application PC/TUS9407659 GENERAL INFORMATION:
                                                                                                                         GENERAL INFORMATION:
APPLICANT: MONTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (610) 270-50s
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 247 amino acid
                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 19406-2799
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
PILING DATE: 09-UUL-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
CORRESPONDENCE ADDRESS:
                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Sutton, Jeffrey REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: SmithKline Beecham Corporation - Corp ADDRESSEE: Intellectual Property STREET: 709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                        80
                                                                                                                                                                                                                                                                                                 61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                       94;
                                                                                                                                                                                                                                                                                                                                                                    1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                King of Prussia
                                                                                                                                                                                                                                                                        PDTVKGRFTISRDNAKNTLYLOMSSLKSEDTAMYHCARGGVRRGYFDVWGAGTTVTVSS 138
                                                                                                                                                                                                                                                                                                                                               EVHLVESGGGLVKPGGSLKLSCAASGFAFSSYDMSWVRQTPEKRLDWVAYISSGGGGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    247 amino acids
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VENTION: Recombinant and Humanized Il-1 beta
VENTION: Antibodies for Treatment of Il-1 Mediated Inflammatory
                                                   P REZ RODR GUEZ,
SIERRA BL ZQUEZ,
TORMO BRAVO, B. I
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Hurle, Mark
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Jonak, Zdenka L.
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                            LOMBARDERO
                                                                                                                           MONTERO CASIMIRO, J. E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (610) 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             270-5090
                                  Anti-CD6 monoclonal antibodies and their uses
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                                                                                                          VALLADARES,
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Pred. No. 1.4e-44;
7; Mismatches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                         18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 247;
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RESULT 8
US-08-553-497A-18
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OTHER INFORMATION: variant of sub-clone for tlA recognizing OTHER INFORMATION: to the variable region of its heavy chains.
                                                                         GENERAL INFORMATION:
APPLICANT: KETTLEB
APPLICANT: BENDIG,
APPLICANT: ANSELL,
                                                                                                                                                     Patent No.
                                                                                                                                                                       Sequence 18, Application US/08553497A
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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FRACYENT TYPE: -N ...
ORIGINAL SOURCE:
TISSUE TYPE: Animal cells.
TISSUE TYPE: NSO " SP 2/0 " CI
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FILING DATE: 17-July-1997
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/CU96/00004
FILING DATE: 18-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
REGISTRATION UNMBER: P-12
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 3:
APPLICANT:
APPLICANT:
APPLICANT:
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HYPOTHETICAL: NO
ANTI-SENSE: NO
TYPE: -N Terminal fragment.
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STREET: V...
CITY: Scarsdale
CTATE: New York
'I S.A.
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LENGTH: 119 Amino acid residues.
TYPE: Amino acid.
STRANDEDNESS: Unknown.
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MEDIUM TYPE: Floppy disk 3.5'' (1.4 MB).
COMPUTER: Compatible PC IBM (80486, 8 M Ram).
                                                         APPLICANT:
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                                                                                                                                                                                                                                                                      61
                                                                                                                                                                                                                                                                                        61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                           94;
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                                                                                                                                                                                                                                                                                                                                              EVQLVESGGGLVKPGGSLKLSCAASGFKFSRYAMSWVRQAPGKRLEWVATISSGGSYIYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sub-clone ior t1A
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MITJANS, FRANSESC
ROSELL, ELISABET
                                                 ANSELL, KEITH H
                                                                                           BENDIG,
                                     ADAN,
                                                                                                                 KETTLEBOROUGH, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
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                                   JAUME
                                                                           KEITH H.
                                                                                               MARY M
                                                                                                                                                                                                                                                                                                                                                                                                                                         76.9%;
79.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 variant of sub-clone ior t1A recognizing human CD6, particula to the variable region of its heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 480; DB 4; Length 119, Pred. No. 2.6e-44; 4; Mismatches 21; Indels
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US-08-553-497A-18
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                                                                                                                                                                                                 Sequence 16, Application US/08579378A Patent No. 6210671
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 530
PRIOR APPLICATION DATA:
RAPPLICATION NUMBER: WO PCT
PRILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION UMBER: EP 941
PILING DATE: 17-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BLASCO, FE
APPLICANT: PIULATS, C
TITLE OF INVENTION: I
                                                                                                                    APPLICANT: Co, Man Sung
TITLE OF INVENTION: Humanized ITITLE OF INVENTION: L-Selectin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 9.
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, VeCURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/553,497A
FILING DATE: 17-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC Com
                                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 703-243-6333
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                ADDRESSEE: Townson, ADDRESSEE: One MarketPlaza,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 75.6%;
Local Similarity 77.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: 33,302
REFERENCE/DOCKET NUMBER: MERCK 1726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                              61 SDSVXGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                          61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92;
                                                                                                                                                                                                                                                                                                                                                                                                                           1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60
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San Francisco
California
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                                                                                                                                                                                                                                                                                                                        PDSVKGRFTISRDNAKNTLYLOMSSLKSEDTAMYYCARLETGDYALDYWGOGTTVTVSS
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                                       Townsend and Townsend and Crew
ne MarketPlaza, Steuart Tower, Suite 2000
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;, JAUME
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ANTIBODIES
                                                                                                                                         Humanized Antibodies Reactive with
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Pred. No. 4.4e-43;
8; Mismatches 19; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 239;
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

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RESULT 10
US-08-326-362-2
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                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application Patent No. 5730981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local 9
                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Bosslet, Klaus
APPLICANT: Seeman, Gerhard
APPLICANT: Dippold, Wolfgang
TITLE OF INVENTION: Monoclonal Anti-Ganglioside Antibody,
TITLE OF INVENTION: Its Preparation and Use as a Tumortherapeutic Agent
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 135 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Liebescheutz, Joe O. REGISTRATION NUMBER: 37,505 REGISTRATION NUMBER: 11,505 REFERENCE/DOCKET NUMBER: 111 TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: EP 95112895.8 FILING DATE: 17-AUG-1995 PRIOR APPLICATION DATA: EP 95114696.8 FILING DATE: 19-SEP-1995 ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/160,074
FILING DATE: 30-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/983,946
FILING DATE: 01-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/579,378A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                    ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
STREET: 1300 I Street, N.W., Suite 700
CITY: Washington
                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 27-DEC-1995 CLASSIFICATION: 424
                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       I: 135 amino acids
   amino acid

XGY: linear
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79.8%;
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Pred. No. 4.1e-43;
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; LENGTH: 119
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-406-535-2
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                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Connie L. Erickson-Miller
APPLICANT: Stephen D. Holmes
APPLICANT: James D. Winkler
TITLE OF INVENTION: Tie2 Antagonist Antibodies
FILE REFERENCE: P50844
CURRENT APPLICATION NUMBER: US/09/406,535
CURRENT FILING DATE: 1999-09-27
EARLIER FILING DATE: 1998-09-28
EARLIER FILING DATE: 1998-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 11
US-09-406-535-2
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                                                                                                                                                   Best Loc
Matches
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Matches
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Patent No. 6376653
                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 118 amino acid
                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/032,863
FILLING DATE: 17-MAR-1993
APPLICATION NUMBER: DE P 42 08 795.3
FILLING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 118 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Einaudi, Carol P. REGISTRATION NUMBER: 32 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
61 ADTVKGRFTISRDNAKNTLFLQMTSLRSEDTAIYYCARGYYGPYYFDYWGQGTALTVSS 119
                     61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 RDSVKGRFTISRDNAKNTLYLQMSSLRSEDTAMYYCARGG-SRYAMDYWGQGTTVTVSS 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 QVQLQQSGGGLVKPGGSLTLSCAASRFTFSTYAMSWVRQTPAKRLEWVAYISSGGASTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60
                                                                                                                                                 h 74.4%;
Similarity 74.8%;
89; Conservative
                                                                       EVQLVESGGGLVKPGGSLKLSCAASGFTFSDYGMHWVRQAPEKGLEWVAYINSGSSTITY 60
                                                                                                         EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                                                                                 Score 464; DB 3; 1
Pred. No. 1.4e-42;
8; Mismatches 22;
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Pred. No. 7.3e-43;
7; Mismatches 19
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                                                                                                                                                                                     Length 119;
                                                                                                                                                     Indels
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US-09-518-737-2

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RESULT 13
US-09-495-880A-11
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                                                                                      US-09-495-880A-11
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Patent No. 6709833
GENERAL INFORMATION:
APPLICANT: PUKUI, YASUHISA
APPLICANT: NAGATA, SATOSHI
APPLICANT: SAIRAI, RYUICHI
APPLICANT: SAITO, NAOAKI
TITLE OF INVENTION: PHONOCLONAL ANTIBODY RECOGNIZING
TITLE OF INVENTION: PHONOCHATIOYLINOSITOL-3,4-DIPHOSPHATE
FILE REFERENCE: 1965/49618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/518,737
CURRENT FILING DATE: 2000-03-03
PRIOR APPLICATION NUMBER: JP 1999-250209
PRIOR FILING DATE: 1999-09-03
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
Query Match
Best Local Similarity
Matches 91; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                  SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11, Application US/09495880A
Patent No. 6667150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 92;
                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/495,880A
CURRENT FILING DATE: 2000-02-01
PRIOR APPLICATION NUMBER: PCT/EP98/04836
PRIOR FILING DATE: 1998-08-03
PRIOR APPLICATION NUMBER: EP 97 11 3319.4
PRIOR PILING DATE: 1997-08-01
NUMBER OF SEQ ID NOS: 50
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: RUDERT, FRITZ
APPLICANT: GE, LIMING
APPLICANT: ILAG, VIC
TITLE OF INVENTION: NOVEL METHOD AND PHAGE FOR THE IDENTIFICATION OF
TITLE OF INVENTION: NUCLEIC ACID SEQUENCES ENCODING MEMBERS OF A MULTIMERIC
TITLE OF INVENTION: (POLY) PEPTIDE COMPLEX
FILE REFERENCE: MORPHO/9
                                                                                                                                                                                        LENGTH: 45
TYPE: PRT
                                                                                         FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: ompA-FLAG-scFv
OTHER INFORMATION: HAG)-gene IIIss encoded by phage vector fhaglA (cir
                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       118
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                                                                                                                                                                                                                456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VSA 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EVQLVESGGDLVKPGGSVKLSCAASGFTFSSYGMSWARQTPDKRLEWVASISSGGSYTYY
73.3%;
ilarity 75.8%;
Conservative
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    8
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Pred. No. 3.4e-42;
Pred. No. 3.7e-42;
Score 457.5; DB .
Pred. No. 3.6e-41
8; Mismatches 2
                                         DB 4;
    20;
                                       Length 456;
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  Indels
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1:
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Gaps
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(circular)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60
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FILE REFERENCE: p1775R1
CURRENT APPLICATION NUMBER: US/09/648,067A
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,018
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/213,822
PRIOR FILING DATE: 2000-06-23
NUMBER OF SEQ ID NOS: 15
SEQ ID NO 15
LENGTH: 119
TYPE: PRT
                                                       ; OTHER INFORMATION: US-09-648-067A-15
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MOLECULE TYPE: protein

PCT-US94-14106-59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII (text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14106
FILING DATE:
CLASSIFICATION:
INFORMATION FOR SEQ ID NO: 59:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 14
PCT-US94-14106-59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 75.6
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 59, Applicat GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15, Application US/09648067A Patent No. 6627196
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Dosages for Treatment with Anti-ErbB2 Antibodies FILE REFERENCE: P1775R1
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Baughman, Shapplicant: Shak Steven
                                                                                            ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 217 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 6
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       170 EVQLVESGGDLVKPGGSLKLSCAASGFSFSSYGMSWVRQTPDKRLEWVATISNGGGYTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               230 PDSVKGRFTISRDNAKNTLYLQMSSLKSEDSAMYYCARRERYDENGFAYWGQGTLVTVSA 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADSVKGRFTISRDNAKNNLYLQMNSLKSEDTAMYYCARDPL----YGSWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EVKLMESGGDLVKPGGSLKLSCAASGFTFSDYYMYWVRQTPEKRLEWVATISDDGTYTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCT-RVKLGTYYFDSWGQGTTLTVSS 119
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                                                                                                                                                                                                                                                                                                                                                                                   Sharon A.
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75.6%;
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72.6%;
71.4%;
                                                                          consensus
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Pred. No. 1.6e-41;
Score 453; DB 4;
Pred. No. 2.1e-41;
                                                                         sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17;
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                   Length 119;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.
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13; Mismatches 21; Indels 0; Gaps 0; 1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60	Job cime :	Search com	문	δ	문	S S	Matches
	Job Elme : 8.85924 secs	Search completed: December 23, 2004, 19:08:01		61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119	1 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY 60	1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60	85; Conservative 13; Mismatches 21; Indels 0; Gaps

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No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-089-500-55
624
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98.4
98.0
77.9
77.1
76.1
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76.1
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Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09C_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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14
US-10-265-713-8

US-10-166-626-8

7 US-10-06-773-4

3 US-10-276-939-3

US-10-276-939-3

US-10-281-479A-61

US-10-286-132A-61

US-10-281-479A-23
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US-10-166-626-18
US-09-764-304-8
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Sequence 18, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 8, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 61, Appli
Sequence 62, Appli
                                                                                                                                                                                                                                                                                                                  Description
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ALIGNMENTS

US-09-764-304-18

Sequence 18, Application US/09764304 Patent No. US20020026036A1 GENERAL INFORMATION: APPLICANT: SHITARA, KENYA

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APPLICANT: HANAI, NOBUC
APPLICANT: HASEJAMA, MAMORU
APPLICANT: HASEJAMA, MAMORU
APPLICANT: KUMANA, VOSHIHISA
APPLICANT: KUMANA, VOSHIHISA
FILE REFERENCE: 249-101
CURRENT APPLICATION PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US/09/255,322
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US/08/408,133
EARLIER APPLICATION NUMBER: US/08/408,133
EARLIER APPLICATION NUMBER: US/08/292,178
EARLIER APPLICATION NUMBER: US/08/292,178
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US/09/47,674
EARLIER APPLICATION NUMBER: US/09/47,674
EARLIER APPLICATION NUMBER: US/09/47,674
EARLIER APPLICATION NUMBER: US/09/47,674
EARLIER APPLICATION NUMBER: US/09/17/47,674
EARLIER APPLICATION NUMBER: US/09/17/47,674
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PRT ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: CDNA KM-641
US-09-764-304-18
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US-10-166-626-18
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US-10-265-713-18
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PRIOR FILING DATE: 1999-01-05
PRIOR PELICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR PELICATION NUMBER: US 08/292,178
PRIOR PELICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US 07/947,674
                                                         Sequence 18, Application US/1016626
Publication No. US20030166876A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18, Application No. US201 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 18
LENGTH: 130
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APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: HASEGÁWA, MÁMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWAMA, YOSHHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SHITARA, KENYA APPLICANT: HANAI, NOBUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: JP 3-238375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                          EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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No. US20030095964A1
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Pred. No. 2.6e-50;
); Mismatches 0;
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Pred. No. 2.6e-50;
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; FEATURE:
; OTHER INFORMATION: cDNA KM-641
US-09-764-304-8
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US-09-764-304-8
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                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US 07/947,674
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-17
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CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR PILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR PRIOR DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR FILING DATE: 1992-09-17
                                                                                                                                       SEQ ID NO 8
LENGTH: 130
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SEQ ID NO 18
LENGTH: 130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
                                                                                                                                                                                            EARLIER APPLICATION NUMBER: JP 3-238375
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
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                                                          LENGTH: 130
TYPE: PRT
ORGANISM: Artificial Sequence
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Local Similarity 100.0%;
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Pred. No. 2.6e-50;
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DB 9; Length 130

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US-10-265-713-8
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US-10-265-713-8
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PRIOR FILING DATE: 1995-01-05
PRIOR PPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR PPLICATION NUMBER: US/947,674
PRIOR APPLICATION NUMBER: US/93-238375
PRIOR PILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
                                     GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                Sequence 8, Application US/10166626 Publication No. US20030166876A1
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Best Local Similarity
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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver.
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 118;
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MIYAJI, HIROMASA
KUWANA, YOSHIHISA
                                       HASEGAWA, MAMORU
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99.2%;
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99.2%;
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Pred. No. 2.2e-49;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 614; DB 14; Length 130; Pred. No. 2.2e-49;
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; LENGTH: 134
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-477-377-3
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CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-51
PRIOR PILING DATE: 1995-05-21
PRIOR PPLICATION NUMBER: US 08/408,133
PRIOR PILING DATE: 1995-03-21
PRIOR PILING DATE: 1995-03-21
PRIOR PILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR PILING DATE: 1992-09-17
PRIOR PILING DATE: 1992-09-17
PRIOR PILING DATE: 1992-09-17
                                                                                                                                        Query Match 78.0
Best Local Similarity 79.3
Matches 96; Conservative
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APPLICANT: Tornetta, Mark
APPLICANT: Tornetta, Mark
APPLICANT: Wattam, Trevor
TITLE OF INVENTION: Useful in Treatment of RANK Ligand Mediated
                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: GP50050
CURRENT APPLICATION NUMBER: US/10/477,377
CURRENT FILING DATE: 2003-11-12
PRIOR APPLICATION NUMBER: PCT/US02/14246
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: 60/292,031
PRIOR FILING DATE: 2001-05-18
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Local Similarity 99.2%;
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61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTY--YFDSWGQGTTLTVS 118
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                                                                          EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                               EVQLVESGGDLVKPGGSLKLSCAASGFTFSRYGMSWVRQTPDKRLEWVATISSGGSYTYY
                                                                                                                                                                 78.0%;
79.3%;
                                                                                                                                      Score 487; DB 17;
Pred. No. 1.4e-37;
6; Mismatches 17;
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Pred. No. 2.2e-49;
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                                                                                                                                             17; Indels
                                                                                                                                                                                       Length 134;
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US-10-276-939-3
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US-10-006-773-4
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                                                                        SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 121
TYPE: PRT
ORGANISM: Murine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/10276939 Publication No. US20030215450A1 GENERAL INFORMATION:
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SEQ ID NO 4
LENGTH: 140
TYPE: PRT
ORGANISM: Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/10006773
Publication No. US20020132983A1
GENERAL INFORMATION:
APPLICANT: Junghans, Richard P.
TITLE OF INVENTION: Antibodies as Chimeric Effector Cell Receptors Against Tumor Anti
Query Match
Best Local Similarity
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Best Local (
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APPLICANT: SWEET, RAYMOND W.
APPLICANT: TAYLOR, ALEXANDER H.
APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: Anti-RANK Ligand Monoclonal Antibodies
TITLE OF INVENTION: Useful in Treatment of RANK Ligand Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 003
CURRENT APPLICATION NUMBER: US/10/006,773
CURRENT FILING DATE: 2001-12-10
                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/276,939
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: PCT/US01/16865
PRIOR FILING DATE: 2001-05-24
PRIOR FILING DATE: 2001-05-24
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PRIOR FILING DATE: 2000-11-30
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/207,628 PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: GP50024
                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
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77.1%;
78.5%;
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Score 481; DB 14; Pred. No. 4.7e-37;
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US-10-281-479A-61
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 61
LENGTH: 119
Type: """
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PRIOR FILING DATE: 2002-06-24
PRIOR PRICINON UNMBER: 60/346,402
PRIOR FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: PCT/US01/14151
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: 60/201,344
PRIOR FILING DATE: 2000-05-02
                                                           Sequence 61, Application US/10275180A Publication No. US20030190687A1 GENERAL INFORMATION:
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  APPLICANT: The UAB Research Foundation APPLICANT: Zhou, Tong APPLICANT: Ichikawa, Kimihisa
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APPLICANT: Buchabaum, Donald J.
TITLE OF INVENTION: COMBINATIONS OF ANTIBODIES SELECTIVE FOR A TUMOR NECROSIS
TITLE OF INVENTION: FACTOR-RELATED APOPTOSIS-INDUCING LIGAND RECEPTOR AND OTHER THER
TITLE OF INVENTION: AGENTS
FILE REFERENCE: 21085-0029U6
CURRENT APPLICATION NUMBER: US/10/281,479A
CURRENT FILING DATE: 2003-01-28
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Local Similarity 79.8%;
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Ichikawa, Kimihisa
Ichikawa, Kimihisa
Kimberly, Robert P.
Koopman, William J.
Oshumi, Jun
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Zhou, Tong
Ichikawa, Kimihisa
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Pred. No. 1.7e-36;
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US-10-286-132A-61
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CURRENT APPLICATION NUMBER: US/10/286,132A
CURRENT FILING DATE: 2003-01-22
PRIOR APPLICATION NUMBER: US 60/346,402
PRIOR FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: PCT/US01/14151
PRIOR FILING DATE: 2001-05-02
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/201,344
PRIOR FILING DATE: 2000-05-02
NUMBER OF SEQ ID NOS: 102
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Best Local :
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SEQ ID NO 61
LENGTH: 119
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Publication No. US20030198637A1
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/275,180A
CURRENT FILING DATE: 2002-10-31
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin version 3.0
SEQ ID NO 61
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APPLICANT: Kimberly, Robert P.
APPLICANT: Koopman, William J.
APPLICANT: LoBuglio, Albert S.
APPLICANT: Buchsbaum, Donald J.
TITLE OF INVENTION: AN ANTIBODY SELECTIVE FOR A TUMOR NECROSIS FACTOR-RELATED
TITLE OF INVENTION: APOPTOSIS-INDUCING LIGAND RECEPTOR AND USES THEREOF
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TITLE OF INVENTION: AN ANTIBODY SELECTIVE FOR A TUMOR NECROSIS FACTOR-RELATED APOPTOS
TITLE OF INVENTION: INDUCING LIGAND RECEPTOR AND USES THEREOF
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ORGANISM: Artificial Sequence
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                                      SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                          EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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PDSVKGRFTISRDNAKNTLYLOMSSLRSEDTAMYYCARRGDSMITTDYWGQGTTLTVSS 119
                                                                                 EVMLVESGGGLVKPGGSLKLSCAASGFTFSSYVMSWVRQTPEKRLEWVATISSGGSYTYY 60
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Pred. No. 1.7e-36;
3; Mismatches 21;
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Pred. No. 1.7e-36;
3; Mismatches 21;
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RESULT 13
US-10-281-479A-23
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PRIOR FILING DATE: 2002-06-24
PRIOR PPLICATION NUMBER: 60/346,402
PRIOR PPLICATION NUMBER: 60/346,402
PRIOR APPLICATION NUMBER: PCT/US01/14151
PRIOR APPLICATION NUMBER: PCT/US01/14151
PRIOR FILING DATE: 2001-05-02
PRIOR PPLICATION NUMBER: 60/201,344
PRIOR FILING DATE: 2000-05-02
NUMBER OF SEQ ID NOS: 102
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Matches
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                                                                                                                                                                                                         APPLICANT: Kimberly, Robert P.
APPLICANT: Koopman, William J.
APPLICANT: LoBuglio, Albert S.
APPLICANT: LoBuglio, Albert S.
APPLICANT: Buchsbaum, Donald J.
TITLE OF INVENTION: AN ANTIBODY SELECTIVE FOR A TUMOR NECROSIS FACTOR-RELATED
TITLE OF INVENTION: APPOPTOSIS-INDUCING LIGAND RECEPTOR AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
CURRENT APPLICATION NUMBER: US/10/286,132A CURRENT FILING DATE: 2003-01-22 PRIOR APPLICATION NUMBER: US 60/346,402 PRIOR FILING DATE: 2001-11-01 PRIOR APPLICATION NUMBER: PCT/US01/14151 PRIOR APPLICATION NUMBER: US 60/201,344 PRIOR FILING DATE: 2001-05-02 PRIOR FILING DATE: 2000-05-02
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APPLICANT: LoBuglio, Albert S.

APPLICANT: LOBuglio, Albert S.

TITLE OF INVENTION: COMBINATIONS OF ANTIBODIES SELECTIVE FOR A TUMOR NECROSIS TITLE OF INVENTION: FACTOR-RELATED APOPTOSIS-INDUCING LIGAND RECEPTOR AND OTHER OF INVENTION: AGENTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 21085.0029U6
CURRENT APPLICATION NUMBER: US/10/281,479A
CURRENT FILING DATE: 2003-01-28
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                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zhou, Tong
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Kimberly, Robert P.
Koopman, William J.
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79.8%;
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Pred. No. 6.8e-36;
3; Mismatches 21;
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CURRENT APPLICATION NUMBER: US/10/275,180A
CURRENT FILING DATE: 2002-10-31
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin version 3.0
SEQ ID NO 23
LENGTH: 464
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:/No. US20030190687A1e =
OTHER INFORMATION: Synthetic Construct
US-10-275-180A-23
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US-10-275-180A-23
; Sequence 23, Application US/10275180A
; Publication No. US20030190687A1
; GENERAL INFORMATION:
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Search completed: December 23, 2004, 19:35:48 Job time: 28.0707 secs
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US-10-286-132A-23
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 23
LENGTH: 462
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Best Local Similarity 79.8%;
Matches 95; Conservative
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APPLICANT: Zhou, Tong
APPLICANT: Zhou, Tong
APPLICANT: Ichikawa, Kimihisa
APPLICANT: Kimberly, Robert P.
APPLICANT: Koopman, William J.
TITLE OF INVENTION: AN WILLIAM J.
FILE REFERENCE: 21085.0029U5
FILE REFERENCE: 21085.0029U5
FILE REFERENCE: 21085.0029U5
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    k; Pred. No. 6.8e-36;
    3; Mismatches 21; Indels

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A;Cross-references: GB:M29252; NID:g195657; PIDN:AAA38388.1; PID:g195658 C;Superfamily: immunoglobulin V region; immunoglobulin homology F;15-98/Domain: immunoglobulin homology <IMM>

Length 119;

Query Match

A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-119 <STU>

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anti-poptide Fab' B13I2 heavy chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 22-Jun-1990 #sequence_revision 22-Jun-1990 #text_change 23-Jul-1999
C;Accession: B34353

R;Stura, E.A.; Stanfield, R.L.; Fieser, T.M.; Balderas, R.S.; Smith, L.R.; Lerner, R. J. Biol. Chem. 264, 15721-15725, 1989
A;Title: Preliminary crystallographic data and primary sequence for anti-peptide Fab' A;Reference number: A34353; MUID:89359424; PMID:2504725
A;Accession: B34353

R.S.; Smith, L.R.; Lerner, R.A.

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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-120 <BOE>
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C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 23-Jul-1999
C;Accession: S5536
C;Accession: S5536
R;Boettger, V.; Boettger, A.; Lane, B.B.; Spruce, B.A.
J. Mol. Biol. 247, 932-946, 1995
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                                                                Cross-references: EMBL:X82589; NID:g854304; PIDN:CAA57925.1; PID:g854305; Superfamily: immunoglobulin V region; immunoglobulin homology; Keywords: heterotetramer; immunoglobulin
                                               ;14-97/Domain:
  Query Match
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                                             immunoglobulin homology <IMM>
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78.2%;
75.5%;
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Pred. No. 1.3e-37;
8; Mismatches 18
  Score 471;
  DB
2
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Length 120,
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EMBO J. 5, 1577-1587, 1986

A; Mitle: Structural and functional implications of a restricted antibody response ha; Mitle: Structural and functional implications of a restricted antibody response ha; Reference number: A91043; MUID:86300658; PMID:2427335

A; Recession: 127887

A; Residues: 1-121 < CATS

A; Experimental source: strain Balb/c

A; Experimental source: strain Balb/c

A; Experimental source: strain Balb/c

A; Note: this sequence was determined from the germline gene
C; Comment: This chain was isolated from a hybridoma protein that binds influenza v
C; Superfamily: immunoglobulin v region; immunoglobulin homology
C; Superfamily: immunoglobulin homology < IMM's
                                                                             A;Title: Comprehensive epitope analysis of monoclonal anti-proenkephalin antibodies usin utations in the variable region genes.

A;Reference number: S55528; MUID:95Z39763; PMID:7536850

A;Accession: S55537

A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-120 <BOE>
A;Residues: 1-120 <BOE>
A;Cross-references: EMBL:X82590; NID:g854306; PIDN:CAA57926.1; PID:g854307
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                               Ig heavy chain V region pe21 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change
C;Accession: S5537
R;Boettger, V.; Boettger, A.; Lane, E.B.; Spruce, B.A.
J. Mol. Biol. 247, 932-946, 1995
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127887
127887
19 heavy chain V region (H37-45) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change C;Accession: 127887
R;Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
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Best Local S
Matches 94
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                    75.2%;
78.3%;
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Score 469; DB 2;
Pred. No. 1.1e-36;
5; Mismatches 19
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Pred. No. 8e-37;
8; Mismatches
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R;Tillman, D.M.; Jou, N.T.; Hill, R.J.; Marion, T.N.
J. Exp. Med. 176, 761-779, 1992
A;Title: Both IgM and IgG anti-DNA antibodies are the pro A;Reference number: PH0971; MUID:92381444; PMID:1512540
A;Accession: PH1007
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-111 <TIL>
A;Experimental source: B cell, strain [NZB x NZW]F1
A;Experimental source: B cell, strain [NZB x NZW]F1
C;Superfamily: immunoglobulin V region; immunoglobulin ho C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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A;Molecule type: DNA
A;Residues: 1-122 <CAT>
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E27888
Ig heavy chain
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C;Species: Mus musculus (house mouse)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PH1007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Experimental source: strain Balb/c
A;Note: this sequence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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C;Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change
C;Accession: E27888
R;Caton, A.J; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
EMBO J. 5, 1577-1587, 1986
EMBO J. 5, 1577-1587, 1986
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Mus musculus (house mouse)
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                     EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
  EVKLVESGGGLVKPGGSLKLSCAASGFTFSSYAMSWVRQTPAKRLEWVAYISSGGGSTYY
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                                                                                                 74.8%;
82.0%;
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                                                                                               Score 467; DB 2;
Pred. No. 1.5e-36;
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Pred. No. 1.1e-36;
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Ig heavy chain V region precursor - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 23-Jul-1999
C;Accession: S09258
R;Hamada, H.; Maezawa, K.; Tsuruo, T.
Nucleic Acids Res. 18, 1900, 1990
A;Title: Nucleotide sequences of the variable regions of a mouse monoclonal ar A;Title: Nucleotide sequences of the variable regions of a mouse monoclonal ar A;Reference number: S09258; MUID:90245594; PMID:2110659
A;Reference number: S09258; MUID:90245594; PMID:2110659
A;Rocession: S09258
A;Molecule type: DNA
A;Residues: 1-138 <HAM's
A;Cross-references: EMBL:X51719; NID:g53207; PIDN:CAA36012.1; PID:g297545
C;Genetics:
A;Introns: 16/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Reywords: heterotectramer; immunoglobulin
F;34-117/Domain: immunoglobulin homology <IMM's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: DNA
A;Residues: 1-119 <CAT>
A;Residues: 1-119 <CAT>
A;Residues: 1-119 <CAT>
A;Experimental source: strain Balb/c
A;Superimental source: strain Balb/c
A;Comment: This equence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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;Species: Mus musculus (house mouse)
;Date: 15-Dec-1988 #sequence_revision
                                                                                                                                                                                        Query Match
Best Local :
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Best Local :
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                       SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                        EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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PDSVKGRFTISRDNAKNNLYLQMSSLRSEDTALYYCARYYRYEAWFASWGQGTLVTVSA 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PDSVKGRFTISRDNAKNTLYLOMSSLKSEDTAMYYCPRGE--TYYDYAMDYWGQGTSVTV
                                                                                EVILVESGGGLVKPGGSLKLSCAASGFTFSSYTMSWVRQTPEKRLEWVATISSGGGNTYY
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76.9%;
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                                                                                                                                                              Score 464; DB 2; 1
Pred. No. 3.7e-36;
7; Mismatches 22;
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Pred. No. 1.8e-36;
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Ig heavy chain V region - mouse C; Species: Mus musculus (house C; Date: 20-Feb-1995 #sequence_x
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F;31-35/Region:
F;50-66/Region:
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Mol. A;Title: Analysis of the structural diversity of monoclonal antibodies to cycle A;Reference number: PH0087; MUID:91042649; PMID:2122240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig heavy chain V region (anti-cyclosporin B) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 15-Jan-1993 #sequence_revision 15-Jan-1993 #text_change
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C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-118 < SCI
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                                                                                             RESULT
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C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change
C;Accession: S55539
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J. Mol. Biol. 247, 932-946, 1995
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                                                                                                                                                                                          SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
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complementarity-determining 1
complementarity-determining 2
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79.08;
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Pred. No. 4.3
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Pred. No. 3.9e-36;
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  20-Feb-1995 #text_change 23-Jul-1999
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Ig heavy chain V region (anti-DNA, 3E12VH) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Dete: 16-Sep-192 #sequence_revision 16-Sep-1992 #text_change 16-Aug-1996
C;Accession: PL0249
C;Accession: PL0249
R;Shlomchik, M; Mascelli, M; Shan, H; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A
J. Exp. Med. 171, 265-297, 1990
A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic
A;Reference number: PL0231; MUID:90111618; PMID:2104919
A;Accession: PL0249
A;Molecule type: mRNA
A;Residues: 1-117 <SHL>
C;Kuywords: heterotetramer; immunoglobulin
F;1-30/Region: framework 1
F;15-98/Domain: immunoglobulin homology <IMM>
F;15-98/Domain: immunoglobulin homology <IMM>
F;15-98/Domain: immunoglobulin homology <IMM>
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R;Iosman, M.; Fasy, T.M.; Novick, K.E.; Monestier, M.
submitted to the EMBL Data Library, February 1992
A;Description: Relationships among antinuclear antibodies from autoimmune
A;Reference number: $20639
A;Accession: $20641
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-118 <LOS>
A;Cross-references: EMBL:X65003; NID:g52602; PIDN:CAA46136.1; PID:g52603
C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: immunoglobulin homology <IMM>
                  Ig heavy chain V region pe22 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 23-Jul-1999
C;Accession: S55538
R;Boettger, V.; Boettger, A.; Lane, E.B.; Spruce, B.A.
J. Mol. Biol. 247, 932-946, 1995
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S55538
    A; Title: Comprehensive
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67-98/Region:
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Pred. No. 5.9e-36;
8; Mismatches 21;
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Pred. No. 4.8e-36;
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Ig heavy chain V region (17/9) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 2
C;Accession: B31790
R;Schulze-Gahmen, U.; Rini, J.M.; Arevalo, J.; Stura, E.A.; Kente
J. Biol. Chem. 263, 17100-17105, 1988
A;Title: Preliminary crystallographic data, primary sequence, and
A;Reference number: A92686; MUID:89034213; PMID:3182835
A;Accession: B31790
A;Molecule type: mRNA
A;Residues: 1-254 <SCH>
C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-120 <BOE>
A;Cross references: EMBL:X82591; NID:g854308; PII
C;Superfamily: immunoglobulin V region; immunogloC;Keywords: heteroctetramer; immunoglobulin
F;14-97/Domain: immunoglobulin homology <IVM>
C;Accession: PH0096
R;Schmitter, D.; Poch, O.; Zeder, G.; Heinrich, G.F.; Kocher, H.P.; Que Mol. Immunol. 27, 1029-1038, 1990
A;Title: Analysis of the structural diversity of monoclonal antibodies A;Reference number: PH0087; MUID:91042649; PMID:2122240
A;Accession: PH0096
A;Molecule type: mRNA
A;Residues: 1-118 <SCH>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology c;Keywords: immunoglobulin homology c;Keywords: immunoglobulin homology c;Keywords: immunoglobulin homology c;Molecule c; immunoglobulin homology c;Molecule c;
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Pred. No. 2.8
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Pred. No. 7.5e-36;
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F;31-35/Region: complementarity-determining
F;50-66/Region: complementarity-determining
F;99-105/Region: complementarity-determining

Query Match Query Match Best Local Similarity 76.5%; Pred. No. 1.5e-35; Matches 91; Conservative 10; Mismatches 15; Indels 3; Gaps 2; Qy 1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60

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          33222222222111111111
109876543210987654321098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                  424
423.5
423.5
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422.5
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seq length: 2000000000
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624
1 EVTLVESGGDFVKPGGSLKV.....KLGTYYFDSWGQGTTLTVSS
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1: uniprot_sprot:*
2: uniprot_trembl:*
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     GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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BAD00406
BAD00534
BAD00491
AAL35877
Q6GMY2
Q6GMY2
HV55 MOUSE
Q9UL71
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Q99KA4
Q920E7
  OGMZU6
CAE45931
BAD00225
BAD00420
BAD00444
Q6PD88
AAH58814
BAD00233
BAD00475
BAD00480
BAD00424
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AAL35865
Q6PI81
AAH41037
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Q91WP5
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Q6mzu6 homo sapien Cae45931 homo sapi Bad00225 camelus d Bad00420 camelus d Bad00444 camelus d Q6pdb8 mus musculu Aah58814 mus musculu Bad00233 camelus d P01783 mus musculu Bad00445 camelus d Bad00440 camelus d Bad00424 camelus d
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Q6gmy2 homo sapien
P18526 mus musculu
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Aal35865 lama glam
Q6pi81 homo sapien
Aah41037 homo sapi
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66.6	66.6	66.7	66.8	66.8	66.8	66.9	66.9	67.0	67.1	67.2	67.2	67.5	67.5
125	125	117	470	470	117	121	118	123	121	465	465	126	122
N	N	N	N	N	,_	N	N	N	N	Ŋ	N	N	N
BAD00509	BAD00439	AAL35875	AAH18747	Q6PJA4	HV54_MOUSE	BAD00525	AAL35882	BAD00234	BAD00459	AAH62335	Q6P6C4	BAD00510	BAD00446
Bad00509 camelus d	Bad00439 camelus d	Aal35875 lama glam	7	Q6pja4 homo sapien	 P18525 mus musculu 	Bad00525 camelus d	Aal35882 lama glam	Bad00234 camelus d	Bad00459 camelus d	Aah62335 homo sapi	Q6p6c4 homo sapien	Bad00510 camelus d	Bad00446 camelus d

ALIGNMENTS

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RESULT 2
CAG34081
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           PAPAP
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Matches 94
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A Briand J.P., Hoebeke J.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ746180; CAG34081.1; -.
R InterPro; IPR003599; Ig.
R InterPro; IPR003599; Ig.
R InterPro; IPR003596; Ig-V.
R InterPro; IPR003596; Ig-V.
R Pfam; PP00047; ig; 2.
R Pfam; PP00047; ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGV; 2.
R SMART; SM00406; IGV; 2.
CAG34081;
CAG34081;
01-JUN-2004
01-JUN-2004
01-JUN-2004
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Q6KB05,
Q5-JUL-2004 (TrEMBLrel. 27, Created)
O5-JUL-2004 (TrEMBLrel. 27, Last sequence update)
S-JUL-2004 (TrEMBLrel. 27, Last annotation update)
ScFv B8E5 protein (Fragment).
Name=scFv B8E5;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-Balb/c;
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                                                                                                                                                                                                                                                                                                                                       120 SS 121
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94; Conservative
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   (TrEMBLrel. 27, (TrEMBLrel. 27, (TrEMBLrel. 27,
                                                                                                                                                      PRELIMINARY;
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77.0%;
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Created)
Last sequence update)
Last annotation update)
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
A Hopkins R.F., Jordan H., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Hopkins R.F., Jordan H., Moore G.J., Rubin G.M., Hong L.,
A Stapleton M., Garcis M.A., Carninci P., Prange C.,
A Stapleton M., Goderland N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Blosak S.A., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Hitlon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
A Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Holton B., Ketteman J.W., Green B.D., Dickson M.C.,
A Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Schein J.E.,
A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
A Hones S. J. Marray M. A.
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Best Local S
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Q99KA4;
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01-JUN-2001 (TremBLrel.
01-MAR-2004 (TremBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peter J.C., Wallukat G., Tugler J., Maurice D., Roegel C
Briand J.P., Hoebeke J.;
"Modulation of the M2 muscarinic receptor activity with
anti-M2 receptor antibody fragments.";
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ746180; CAG34081.1; -.
"Generation and initial analysis of more than 15,000 and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002). [2]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Mammary tumor metastatized to lung. Tumor arose MEDLINE=22388257; PubMed=12477932; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
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Mammalia; Eutheria;
                                                                                                      Jones S.J., Marra M.A.;
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Rodentia;
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Rodentia;
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77.0%;
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Pred.
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annotation update)
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No. 4.3e-41;
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                                                                         full-length
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Best Local S
Matches 94
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Best Local S
Matches 93
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InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; C1-eet; 2.
Pfam; PP00047; Ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q920E7;
Q920E7;
01-DEC-2001
01-DEC-2001
01-OCT-2003
                                                                                                                                                                                                                              PROSITE; PS50835; IG LIKE; 1.
NON_TER 1 1 1
                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AF307937; AAL09421.1; PIR; C25913; C25913. HSSP; P01783; 1IGC.
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Submitted (SEP-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN
SEQUENCE 487 AA; 52554 MW; 7DG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro;
InterPro;
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HSSP; P01810; 2FBJ.
InterPro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                                     Pfam; PF00047; 19;
SMART; SM00406; IG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pterin-mimicking
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                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (MAR-2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=Czech
                                                                                                                                                                                                                                                                                                                                                                    nterPro; IPR007110;
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SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTY--YFDSWGQGTTLTVS
                                                                                EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                             IPR003596;
                                                                                                                                                                                                                 119 AA;
                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TIRMBLrel. 19, Created)
(TIRMBLrel. 19, Last sequence update)
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Cking anti-idiotope heavy chain variak
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata;
Rodentia;
                                                                                                                                                                                                                   13025 MW;
                                                                                                                                                                                                                                                                                                                                         Ig-like.
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75.0%;
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77.7%;
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to the EMBL/GenBank/DDBJ
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                                                                                                                              6;
                                                                                                                              Score 463; DB 2;
Pred. No. 7.8e-41;
6; Mismatches 17
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Pred. No. 1e-
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                                                                                                                                                                                                                   F6E904044381CA7C CRC64;
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7DC8E96DB333077B CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        O., Cotton
J databases.
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; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     region
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                                                                                                                                 4.
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RC Expression driven by an MMTV-LTR enhancer;

RX MEDLINE-22388257; PubMed=12477932;

RX MEDLINE-22388257; PubMed=12477932;

RX Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,

RA Strausberg R.L., Peingold E.A., Grouse L.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M.J., Uddin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Richarde S., McGwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., McGwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Vilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Vilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Fahey J., Helton E., Kettenan M., Madan A., Rodrigues S., Sanchez A.,

RA Hhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalka U., Smailus D.E., Schnerch A., Schein J.E.,

RA Jones S.J.. Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     문
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                                                                                                                                                                     Query Match
Best Local S
Matches 88
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InterPro; IPRO03597; Ig_cl.
InterPro; IPRO03596; Ig_wHC.
InterPro; IPRO03596; Ig_v.
Pfam; PF07654; Cl-sec; 2.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_MHC; UNKNO; SEQUENCE 486 AA; 52681 MW; 4
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TISSUE=Mammary tumor metastatized
TYPE TO A THE TOTAL THE T
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"Generation and initial analysis of more than 15,000
"Generation and initial analysis of more than 15,000
and mouse cDNA sequences.";
proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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Mammalia; Eutheria;
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01-MAR-2004 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BC010324; AAH10324.1;
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20
                                                                                                                                                                                                              Similarity
EVHLVESGGGLVKPGGSLKLSCVVSGFSFTSYDMSWVRQTPERRLEWVAAITSGGN-TYY
                                                                                EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata;
Rodentia;
                                                                                                                                                                                                              72.9%;
71.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19,
19,
                                                                                                                                                                                                                                                                                                                                                                                        UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
                                                                                                                                                                 Score 455; DB 2;
Pred. No. 2.7e-39;
4; Mismatches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                       4FEF835125DA870B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      enhancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to lung. MMTV-LTR/Wnt1 model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  486
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                                                                                                                                                                                                                                    20
                                                                                                                                                                     16;
                                                                                                                                                                                                                                                 Length 486;
                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     full-length human
                                                                                                                                                                 6
                                                                                                                                                                 Gaps
78
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                                                                                                                                                                                                     Query Match
Best Local S
Matches 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WEDLINE-22388257; Pubmede12477932;

XX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

XA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

XA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

XA Hopkins R.F., Jordan H., Moore T., Max G.I., Wang J., Hsieh F.,

XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

XA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

XA Stapleton M., Joares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

XA Stapleton M., Joares M.B., Bonaldo M.F., Casavant T.L., Prange C.,

XA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

XA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

XA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XA Holards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

XA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

XA Whiting M., McMan A., Young A.C., Shevchenko Y., Bouffard G.G.,

XA Holes B. M. T., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

XONES S. M. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

XONES S. M. S., Schein J.E.,

XO
                                                                                                                                                                                                                                                                                                                                  Submitted (SEP-2001) to the EMB Submitted (SEP-2001) to the EMB EMBL; BC013656; AAH13656.1; -. HSS; P01789; IMCP.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; C1-set; 2.
Pfam; PP07654; C1-set; 2.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_MIC; UNKNO SEQUENCE 479 AA; 51603 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q91WP5;
Q91WP5;
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Generation and initial analysis of more than 15,000 and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
Eukaryota; Metazoa; (
Mammalia; Eutheria; l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-2001 (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strausberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
STRAIN=FVB/N; TISSUE=Colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=FVB/N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     116
   13
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19
                                                                    20
                                                                                                                                                                                                                                            Similarity
                                                                                                               EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TVSS 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TVSS 119
                                                                    EVQLVESGGGLVKPGGSLKVSCAASGLTFSNYAMSWVRQSPEKRLEWVAAINSNGGNTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
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                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chordata;
Rodentia;
                                                                                                                                                                                                                                     72.1%;
73.9%;
                                                                                                                                                                                                                                                                                                                                             UNKNOWN 2.
MW; ECB2D0877748584F CRC64;
                                                                                                                                                                                                        9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                     Score 450; DB 2;
Pred. No. 8.9e-39;
9; Mismatches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99:16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            479
                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        full-length human
                                                                                                                                                                                                                                                                             479
                                                                                                                                                                                                     4;
                                                                                                                                                                                                     Gaps
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                                                            Query Match
Best Local S
Matches 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A Straubberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Prange C.,

A Stapleton M., J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Biosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.J., Hullyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Villalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Hitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

B Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

A Jones S. J. Mayra M.A. .
                                                                                                                                                                                                                                            MGD; MGI:2144967, AU044919.
InterPro; IPR000345; CytC heme_BS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig-c1.
InterPro; IPR003597; Ig-MHC.
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003596; Ig-MHC.
IfterPro; IPR003596; Ig-MHC.
                                                                                                                                                                                                                                                                                                                                                                      submitted (JUL-2001) to the EMBL; BC010327; AAH10327.1; PIR; S68213; S68213. HSSP; P01783; 1IGC. MGD. WGT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Warzywinen.
Jones S.J., Marra M.A.;
"Generation and initial analysis
and mouse cDNA sequences.";
""" Acad. Sci. U.S.A. 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "LISSUE=Mammary tumor metastatized to Expression driven by an MMTV-LTR enha MEDLINE=22388257; PubMed=12477932; Stransherr b T =-1--13 - 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q91Z05;
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Mammary tumor 
Expression driven by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-2001
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q91Z05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=CZECH II;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=CZECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=AU044919;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-DEC-2001 (TrEMBLrel. 19, Created)
1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                           PF00047; ig; 1.
; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80
                                                              86;
                                                                             Similarity
EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                 PS00190; CYTOCHROME_C; UNKNOWN_1.
PS50835; IG_LIKE; 4.
PS00290, IG_MHC; UNKNOWN 1.
473 AA; 51946 MW; CF625F00893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence AU044919
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chordata;
Rodentia;
                                                                             70.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           metastatized to lung. MMTV-LTR/Wnt1
an MMTV-LTR enhancer.;
                                                            10;
                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL/GenBank/DDBJ databases.
                                                          Score 439.5; DB 2
Pred. No. 1.1e-37;
0; Mismatches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99:16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                     CF625F008932AF12 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enhancer.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0£
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         more than 15,000
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                                                                                              473;
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RESULT 9
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02-MAR-2004
02-MAR-2004
02-MAR-2004
                                    NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BAD00534;
02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
10-MAR-2004) (TremBLrel. 27,
                                                                                        EMBL;
                                                                                                                         Honda T., Akahori Y., Kurosawa Y., "Libraries of heavy-chain antibodies gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NON TER
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gamma3 in vivo repertoires."
Submitted (SEP-2002) to the I
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"Libraries of heavy-chain antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
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      SEQUENCE
                                                                                                                                                                                                                                                     TISSUE=Spleen;
                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Camelus dromedarius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunoglobulin
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                                                                                                  AB092172;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
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124
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                                                                                                  BAD00534.1;
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13856 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                  (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
      13421
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      W.
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Last annotation update)
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Last annotation update)
                                                                                                                               EMBL/GenBank/DDBJ
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Pred. No. 3
      B31996777375A27E CRC64
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RESULT 11
AAL35877
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AC AAL35
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DT 02-M
DT 02-M
DT 02-M
DT 02-M
DT 02-M
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OC Lama
OC Lama
OC Mamma
OC MCBLI
RN [1]
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Best Local S
Matches 81
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Best Local
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AALJ5877 PRELIMINARY; PRT; 117 AA.
AALJ5877;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunglobulin heavy chain variable domain (Fragment).
Lama glama (Llama).
Enkeryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Lama.
NCBI TaxID=9844;
[1]
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"Libraries of heavy-chain antibodies reflecting camel gamma2
gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AB092129; BAD00491.1; -.
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Mammalia; Eutheria;
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125 AA;
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ilarity 67.2%;
Conservative 1
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13593 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69.7%;
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Last
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Pred. No. 1.1e-37;
LB; Mismatches 19;
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Pred. No. 7.3e-38;
7; Mismatches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F1637892B028E48C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      region
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annotation update)
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RESULT RECEIVED RECEI
              SEQUENCE FROM N.A.

TISSUB-Primary B-Cells;

Strausberg R.;

Submitted (JUN-2004) to the EMBL

R EMBL; BC073758; AAH73758.1; -.

JR InterPro; IPR003599; Ig.

JR InterPro; IPR003597; Ig-11ke.

DR InterPro; IPR003597; Ig-21.

DR InterPro; IPR003596; Ig_MIC.

TR InterPro; IPR003596; Ig_v.
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WREDLINE=2238857; PubMed=12477932;

A Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschal S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Robask S.A., WcEwan P.J., McKerman K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Hhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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Q6GMY2;
05-JUL-2004 (1
05-JUL-2004 (1
05-JUL-2004 (1
Hypothetical E
  InterPro; IPRO(
Pfam; PF07654;
Pfam; PF00047;
                                                                                                                                                                                                                                                                                                                                                                            Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 and mouse cDNA sequences.":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON_TER
NON_TER
SEQUENCE
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Mammalia; Eutheria;
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J. Immunol. Methods 263:97-109(2002).
EMBL, AF442946; AAL35877.1; -.
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MEDLINE=22007448; PubMed=12009207;
Tanha J., Dubuc G., Hirama T., Nar
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Primates;
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Last sequence update)
Last annotation updat
                                                                                                                                                                                                                    EMBL/GenBank/DDBJ
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Pred. No. 1.1e-37;
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                                                                                                                                                                                                                                                                                                                                                     99:16899-16903 (2002)
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                                                                      Query Match
Best Local S
Matches 82
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Best Local S
Matches 86
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NON TER
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SMART; SM00407; IGC1; 4.
SMART; SM00406; IGC; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS90290; IG_MHC; UNKNOWN_3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE SERVICE MEDILINE=89279149; PubMed=2499654; MEDLINE=89279149; PubMed=2499654; Lebecque V.V., Lebecque in mutation in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOUSE
                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00047; ig; 1. SMART; SM00406; IGv;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Early onset of somatic mutation primary immune response.";
J. EXP. Med. 169:2007-2019(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1990
15-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                             PIR; JT0502; HVMS34.
HSSP; P01783; 1IGC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig heavy chain V reg
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                              SIGNAL
                                                                                                                                                                                                                                                                                                                                                   PROSITE
                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
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                                                                        82;
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; PS50835; lo__
'--1in V region; f
                                                                                          Similarity
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                      EVTLVESGGDFVKFGGSLKVSCAASGFAFSHYAMSWVRQTPAKKLEWVAYISSGGSGTYY
EVQLVESGGGLVKPGGSLKLSCAASGFAFSSYDMSWVRQTPEKRLEWVAYISSGGGSTYY
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(Rel. 16, Last sequence update)
(Rel. 38, Last annotation update)
ain V region 345 precursor.
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46955021
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Rodentia;
                                                                                      69.1%;
83.7%;
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64.2%;
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                                                                                                                                                                                                                                                                                                                              Signal.
                                                                                                                                             MW;
                                                                                                                                                                            Ig heavy chain V region 345.
Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
By similarity.
                                                                   Score 431; DB
Pred. No. 1.8e
5; Mismatches
                                                                    5
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
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                                                                                                                                             49380E4627ACA99A CRC64;
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                                                                   DB 1; I
L.8e-37;
hes 11;
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VH genes
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; Murinae; Mus
                                                                                                         117;
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Best Local S
Matches 82
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Q91XE1;
01-DEC-2001
01-DEC-2001
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Q9UL71;
01-MAY-2000
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NON_TER
SEQUENCE
                 SEQUENCE FROM N.A.
STRAIL=FVB/N; TISSUE=Colon;
MEDLINE=22388257; PubMed=12477932;
Strausberg R.L., Feingold B.A., Gr
Klausner R.D., Collins F.S., Wagne
                                                                                                                                                                                                                                         01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
                                                                                                                                                                                                      ign-vJ558 protein
Name=Igh-vJ558;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Euther
NCBI_TaxID=9606;
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                                                                                                                 NCBI_TaxID=10090;
                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clin. Immunol. Immunopathol. EMBL; AF035043; AAD56279.1;
                                                                                                                                                                                       Mus musculus (Mouse)
                                                                                                                                                                                                                            Igh-VJ558
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tetus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYY--FDSWGQGTTLTVS
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IPR003596; Ig_v.
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121 AA;
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(TrEMBLrel.
(TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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                                                                                                                                                  Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13154 MW;
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13,
25,
d B.A., Gir
S., Wagner L.,
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Last
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Pred. No. 2.4e
14; Mismatches
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Last annotation update)
                                                                                                                                                                                                                                                                                  Created)
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                                                                                                                                                  Craniata; Vertebrata; | Sciurognathi; Muridae;
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Catarrhini;
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                                        Grouse L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ä
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Shenmen
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l; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 121;
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                                        J.G.,
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RA Brownstein M.J., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
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RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Butterfield Y.S.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
"Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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Matches 85
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Interpro; IPR00359; Ig_cl.
Interpro; IPR003595; Ig_wHC.
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Pfam; PF07654; Cl-set; 2.
Pfam; PF00047; ig; 1.
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SMART; SM00406; IGv; 1.
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SEQUENCE 480 AA; 51936 MW; 20B92341
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Strausberg R.;
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; BC010798; AAH10798.1; -.
HSSP; P01789; IMCP.
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ALIGNMENTS

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Query Match Best Local Similarity Matches 108; Conservative

100.0%; sc. 100.0%; Pr

Score 572; DB 4; I Pred. No. 3.3e-39;

Length 108; Indels

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Sequence 108

A,

The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention

Claim 11; Page 174-175; 183pp; Japanese.

activity

in cancer.

WPI; 2001-266143/27.

New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Drugs containing genetically-modified antibody against ganglioside GD3, its fragment, immunocompetent cell activators or/and antitumor agents i combination, applicable in treating malignant tumor like melanoma.
                                                               AAB01628 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ganglioside GD3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified ganglioside GD3 antibody associated protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                               108;
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                                                            protein; 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 572; DB 6;
Pred. No. 3.3e-39;
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RESULT 4
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                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                The present sequence is a murine immunoglobulin light chain variable region from plasmid KM-641. The coding sequence was used in the creation of an expression vector, along with the sequence for a human antibody, to produce humanised chimaeric antibodies, which can be used to treat cancer. Humanised chimaeric antibodies are more effective than mouse antibodies as they do not provoke a reaction in the human and side effects, such as the formation of anti-mouse immunoglobulin antibody and the rapid half-life of the immunoglobulins, do not occur
Ganglioside; GD3; complementarity determining region; CDR; antibody;
                                                                                                                                                                                                                                                                                                                            Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New humanized chimera antibody KM-871 useful for treating cancer, comprises variable region of mouse monoclonal antibody, reactive
                         Ganglioside GD3 specific antibody related protein SEQ ID NO:
                                                    03-JUL-2001
                                                                                                      AAB81978 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 14; Page 28-29; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ganglioside and human antibody constant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-SEP-1991;
18-SEP-1992;
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                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                       RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                                                                                                              RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                                                                                                                                                                                                        DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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                                                   (first entry)
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92EP-00116026.
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/label=_signal_peptide
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/label= mature_immunoglobulin_light_chain_V_region
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                                                                                                      protein;
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Best Local &
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                                                                                                                                    Mus musculus
                                                                                                                                                       Ganglioside
                                                                                                                                                                             Modified
                                                                                                                                                                                                 04-FEB-2003
                                                                                                                                                                                                                                        ABU11003 standard; protein; 128
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 140; 183pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therap of e.g. tumors, with low antigenicity, little side effects but potent
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06-APR-2000;
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          Shitara
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                              (KYOW)
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                                                                                                                                                                            ganglioside
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                                                                                                                                                        GD3;
          Niwa
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                                                  2001JP-00097483
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                                                                                                                                                       anti-ganglioside
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                                                                                                                                                                            GD3 antibody associated protein
          Kanazawa
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                                                                                                                                                                                                                                                                                                                                                                              Score 572; DB 4;
Pred. No. 3.9e-39;
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          Asada
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Best I
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prior art
human anti
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                      This immunoglobulin region was isolated from pKM641LA2. A methionine codon, presumably the initiation codon ATG, was found in the vicinity the 5' terminus and the sequence has a full length leader sequence. The chimeric human antibodies are useful in the treatment of cancer, especially that which is of neural ectodermal origin. In contrast to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drugs containing genetically-modified antibody against ganglioside GD3 its fragment, immunocompetent cell activators or/and antitumor agents combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                    Chimeric human
                                                                                                                                                                                                   Nakamura K,
                                                                                                                                                                                                                                                         07-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                            antibody; nucleotide; genomic;
light chain; amino acid.
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                                                                                                         Example
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                                                                                                                                                                          1999-468416/39
 art constructs based on mouse monoclonal antibodies, antibodies do not cause anti-mouse immunoglobulin pr
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                                                                                                        1; Col 99; 188pp;
                                                                                                                                                               AAX99481.
                                                                                                                                                                                                                               KYOWA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                    immunoglobulin
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                                                                                                                                                                                                   Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                               HAKKO
                                                                                                                                  antibody expression
                                                                                                                                                                                                                                                          93US-00116778
                                                                                                                                                                                                                                                                                   9508-00483528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry.
                                                                                                                                                                                                                               KOGYO
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                                                                                                                                                                                                     Kuwana
                                                                                                         English.
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                          hypervariable region; chimeric;
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                                                                                                                                                                                                     Hasegawa
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. No. 3.9e-39;
smatches 0;
                                                                                                                                    vectors
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tion. The
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
           Query Match
Best Local Similarity
                                                             variable regions respectively. The DNA sequences encoding these proteins were used in the construction of humanised chimeric antibody expression vectors. In these humanised antibodies none of the amino acids of the nor human animal Ab variable region have been changed. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chimeric human antibodies have a prolonged half-life and a reduced frequency of adverse effects when compared to mouse monoclonal antibodies
                                                                                                                                                          Humanised chimeric antibody prodn. against ganglioside {\tt GD3} -cancers, such as melanoma, neuroblastoma, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
12-JUL-1993
                                            Sequence
                                                                                                                 The sequences given in AAR33256-57 represent rat heavy and light chain
                                                                                                                                         Claim 6;
                                                                                                                                                                                             N-PSDB;
                                                                                                                                                                                                                           Shitara K,
                                                                                                                                                                                                                                                                     18-SEP-1991;
                                                                                                                                                                                                                                                                                           18-SEP-1992;
                                                                                                                                                                                                                                                                                                                24-MAR-1993
                                                                                                                                                                                                                                                                                                                                    EP533199-A2
                                                                                                                                                                                                                                                                                                                                                                   Protein
                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                          Promoter; variable; region; rat; immunoglobulin; heavy; H; chain; humanised; chimeric; antibody; expression vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat immunoglobulin L chain varible region of pKM641LA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR33257
                                                                                                                                                                                                                                                (KYOW ) KYOWA HAKKO KOGYO CO LTD
                                                                                                                                                                                                      1993-095510/12.
   107;
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                                                                                                                                      Page 30-31; 63pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                            128
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                                                                                                                                                                                                                           Hanai N,
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  Conservative
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                            note= "Signal peptide"
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99.1%;
          98.3%;
99.1%;
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Score 562; DB
Pred. No. 2.5e
0; Mismatches
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Pred. No. 1.7e-38;
0; Mismatches 1;
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                                                                                                                                                                                                                           Miyaji H,
           DB 2;
.5e-38;
                                                                                                                                                                                                                            Kuwana
                    Length 128;
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RESULT 8
AAR53340
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CONB
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                                                                                                                                                                                                Query Match
Best Local Simi
Matches 106;
                                                                                                                                                                                                                                                                                                                                              Example 2 describes the construction of the vector pChi641HAl for chimeric human antibody H chain expression. mRNA from mouse anti-CD3 monoclomal Ab KM641-producing cells was isolated and KM641 H and L chain cDNA slabtated. The base sequences of the Ig variable regions in KM641 H chain cDNA (pKM641HA3) and KM641 L chain cDNA (pKM641LA2) are given in AAQ45438-39. A KM641-derived chimeric human Ab H chain expression vector was constructed by joining the H chain variable region gene from pKM641HA3 to a vector for chimeric human Ab H chain expression using the synthetic DNAs given in AAQ63439 and AAQ63440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibody; Ab; ganglioside GM2; chimera; chimeric antibody; expression vector; heavy; light; chain; hypervariable region; CDR; constant region; hybridoma; Ig; immunoglobulin; promoter; enhancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Humanised antibody specific cytocidal effect on cancers
                                                                                                                                                                                                                                                                                                    Sequence 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 115-116; 191pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (КУОМ ) КУОМА НАККО КОБУО КК.
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DB; AAQ45438.
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                                                                                                 RFSGGGSGTDYSLTISNLEPGDIATYFCHQYSKLPWTSGGGTKLEIKR
                                            RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
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/label= sig_peptide
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                                                                                                                                                                                                                          97.4%;
98.1%;
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                                                                                                                                                                                                Score 557; DB 2; I
Pred. No. 6.5e-38;
Pred. No. 6.5e-38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            melanoma, neuroblastoma
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                                                                                                                                                                                                                                                  Length 128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used for producing a roblastoma and glioma.
                                                                                                                                                                                                     Indels
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RESULT 10
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AC AAR12
AC AAR12
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DT 15-AU
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Best Local S
Matches 98
                                                                                                                                                                                                                                                                                                                                          The present sequence is the light chain variable region of the anti-human Fas ligand monoclonal antibody (MAb) NOK-1. NOK-1 is produced by the hybridoma NOK-1 (FERM BP-5044), which was prepared by immunising mice with transformed human Fas ligand expressing COS cells, and fusing spleen cells isolated from the mice with myeloma P3x63Ag8.653 (ATCC CRL-1580) cells. The MAb recognises the human Fas ligand on the cell surface or in solution, and can be used to inhibit the application of the cell surface Fas ligand/Fas reaction. The MAb can also be used for a Fas ligand assay in biological samples (e.g. human blood), especially for disease catalogues. e.g. hepatitis, infectious mononucleosis and systemic lupus
                                                                                                                                                                                                                                                                                                      Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Variable region; light chain; human; Fas ligand; monoclonal; antibody; NOK-1; hybridoma; inhibition; apoptosis; assay; diagnosis; disease; hepatitis; infectious mononucleosis; systemic lupus erythematosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monoclonal antibody specifically recognising the Fas ligand -
the detection of Fas ligands either on cell surface or in solu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kayagaki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAR-1995;
27-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW00834 standard; protein;
25-MAR-2003
15-AUG-1991
                                         AAR12359;
                                                                   AAR12359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                                                                                      μ.
                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                            DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKDDGTVKLLIFYSSNLHSGVPS
                                                                                                                                                      RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                   standard;
                                                                                                                                       RESGSGSGTD
                                                                                                                                                                                          DIQMTQTTSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yagita H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
(revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95JP-00087420.
95JP-00303492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96WO-JP000734.
                                                                   protein;
                                                                                                                                  91.1%;
90.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Okumura K,
                                                                   127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                108
                                                                                                                                                                                                                                              Score 521; DB 2;
Pred. No. 4.7e-35;
3; Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                   ζ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nakata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibody NOK-1.
                                                                                                                                                                                                                                                                         Length 108
                                                                                                                                                                                                                                                 Indels
                                                                                                                                       108
                                                                                                                                                                108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              solution.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful
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                                                                                                                                                                                             60
                                                                                                                                                                                                                      60
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밁 Ś 밁 S

Mus musculus

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RESULT 11
AAW04177
ID AAW04
XX AAW04
XX 19-MA
XX Uaria
XX Varia
XX Varia
XX Varia
XX Varia
XX NOK-11
KW NOK-11
KW Varia
XX Varia
XX MAB MM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the light (kappa) - chain variable (V) region of a mouse monoclonal antibody (MAb), ICI1, and is specific for an HIV-1 viral antigen. It is used in the construction of a chimeric MAb comprising heavy and light chains having murine V regions and human C regions. The chimeric MAbs are more effective than murine MAb ICI1 since they have an increased compatibility in humans. The heavy and light chain V-regions are joined by manipulating their respective joining (J) regions, to generate restriction enzyme recognition sites. The chimeric MAbs can be used as immunoconjugates, in association with e.g. toxins for HIV treatment. They can also be used in diagnosis of HIV. See also AAQ12056-60 and AAQ12056-63. (Updated on 25-MAR-2003 to correct pA field.)
(Updated on 25-MAR-2003 to correct PI field.) (Updated on 25-MAR-2003 to
                                                   Variable region; light chain; human; Fas ligand; monoclonal; NOK-1; hybridoma; inhibition; apoptosis; assay; diagnosis; dihepatitis; infectious mononucleosis; systemic lupus erythemat
                                                                                                                                 Variant variable light chain
                                                                                                                                                                       19-MAY-1997
                                                                                                                                                                                                          AAW04177;
                                                                                                                                                                                                                                                AAW04177 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 127 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New chimeric mouse-human antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1991-178105/24.
N-PSDB; AAQ12061.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (XOMA )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-1991.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chiméric antibodies;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Light (kappa) chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Better MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9107493-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antigen from sample.
                                                                                                                                                                                                                                                                                                                                                                                 61
                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ш
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                XOMA CORP.
GREEN CROSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         DIQMTQTASSLEASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                  RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIK 107
                                                                                                                                                                                                                                                                                                                                                                                                                       DIQMTQTTSSLSASLGDRVTISCSASQGISNYLNWYQQKPDGTVKLLIYYTSSLHSGVPS
                                                                                                                                                                                                                                                                                                                                            RESGSGSGTDYSLTISNLEPEDIATYYCQQYSKLPWTFGGGTKLEIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fig 13; 107pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Horwitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                     (first
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                                                                                                                                                                                                                                                protein;
                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      variable region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90.9%;
91.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunoconjugates; HIV; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ghoshdasti P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 520; DB 2;
Pred. No. 6.6e-35;
4; Mismatches 5
                                                                                                                                 of Fas
                                                                                                                               ligand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used
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                                                                                                                                   antibody NOK-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ដូ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detect,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 127;
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RESULT 12
AAW16620
ID AAW16
XX AAW16
XX 14-JA
DT 14-JA
XX Liight
KW monoc
KW hepatt
KW gluta
XX hepatt
XX 
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a variant light chain variable region of the anti-
human Fas ligand monoclonal antibody (MAb) NOK-1. NOK-1 is produced by
the hybridoma NOK-1 (FERM BP-5044), which was prepared by immunising mice
with transformed human Fas ligand expressing COS cells, and fusing spleen
cells isolated from the mice with myeloma P3x63Ag8.653 (ATCC CRL-1580)
cells. The MAb recognises the human Fas ligand on the cell surface or in
solution, and can be used to inhibit the apoptosis inducing cell surface
Fas ligand/Fas reaction. The MAb can also be used for a Fas ligand assay
in biological samples (e.g. human blood), especially for disease
diagnosis, e.g. hepatitis, infectious mononucleosis and systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAR-1995;
27-OCT-1995;
     27-OCT-1995;
                                                       24-OCT-1996;
                                                                                                            01-MAY-1997.
                                                                                                                                                               WO9715326-A1
                                                                                                                                                                                                                                                                                               monoclonal antibody; MAb; hybridoma; treatment; hepatitis; hepatitis B virus; HBV; hepatitis C virus; HCV; apoptosis; liver cell;
                                                                                                                                                                                                                                                                                                                                                       Light chain; variable region; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                         Anti-human FasL antibody (NOK1) light chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW16620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibody specifically recognising the Fas ligand - useful the detection of Fas ligands either on cell surface or in solution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kayagaki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-SEP-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
97; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUMITOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DIQMTQSPSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESGSGSGTDYSLTISNLEPEDIATYECQQYSEFPWTFGGGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 80-81; 133pp; Japanese
                                                                                                                                                                                                                                                                         oxaloacetate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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95JP-00303492.
     95JP-00303491
                                                       96WO-JP003089.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90.2%;
                                                                                                                                                                                                                                                                      pyruvate transaminase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.
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Pred. No. 1.2e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ဗ
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                                                                                                                                                                                                                                                                                                                                                    murine; human; Fas ligand;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                    FasL;
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AAB81998
XX
AC AAB8
XX
AC AAB8
XX
DT 03-(
XX
Gan;
KW Gan;
KW Gan;
KW Can;
XX
OS Syn;
O
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expressed by the hybridoma NOK1 (FERM BP-5044). The MAD can be used in the preparation of a composition for the effective oral or parenteral treatment of hepatitis, including hepatitis caused by hepatitis B or C virus. The composition controls apoptosis in liver cells caused by the binding of FasL to Fas expressing liver cells, and improves liver function by improving blood glutamate oxaloacetate and pyruvate transaminase levels. The composition is given in a dosage of 0.0001-1000, preferably 0.01-600 mg/day. Spleen cells from mice immunised with FasL expressing COS cells were fused with mouse myeloma cells to produce hybridomas. The hybridomas were screened for anti-FasL activity, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is the light chain variable region of the anti-human Fas ligand (FasL) monoclonal antibody (MAD) NOK1, we expressed by the hybridoma NOK1 (FERM BP-5044). The MAD can be
                        WPI; 2001-266143/27.
N-PSDB; AAF86912.
                                                                                                                                                                                                         30-SEP-1999;
06-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ganglioside;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ganglioside GD3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB81998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB81998 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 6; Page 30-31; Slpp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-human Fas Ligand antibody
liver cells and improves liver
                                                                                                   Hanai N,
                                                                                                                                                                                                                                                                                       29-SEP-2000; 2000WO-JP006774
                                                                                                                                                                                                                                                                                                                                                                                             WO200123432-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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97; Conserv
                                                                                                                                                         KYOWA HAKKO
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                                                                                                   Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kayagaki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GD3;
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                                                                                                                                                                                                                                    99JP-00278291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           specific antibody related protein #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        complementarity determining
                                                                                                                                                           KOGYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90.2%;
                                                                                                        Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yagita
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Pred. No. 1.2e-34;
4; Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ħ,
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                                                                                                        <u>,</u> 7
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                                                                                                        Niwa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 108;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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 positive disease.
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Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
             New fusion protein comprising functionally linked components of an au CD7 antibody or its fragment and an immunotoxin, useful for treating positive acute T-cell and/or myeloid leukemias, or Graft-versus-host
                                                                                                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pseudomonas
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                     11-DEC-2002; 2002WU-EP014064
                                                                                                                                                                                                                                                                                     WO2003051926-A2.
                                                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-CD7 antibody TH-69 scFv-Pseudomonas exotoxin A fusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 128 AA;
                                                                                                                                                                                        14-DEC-2001; 2001US-0339422P.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CD7; antibody; TH-69; scFv; exotoxin A; immunotoxin; cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR62591 standard; protein; 650
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                                                                                                                                                          (UYER-) UNIV ERLANGEN-NUERNBERG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunosuppressive;
                                                                                          2003-523519/49.
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                                                                                                                           Gramatzki
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286. .646
                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                 /label= ETA
                                                                                                                                                                                                                                                                                                                                                                             label= scFv
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                                                                                                                        Peipp M;
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Pred. No. 1.4e-34;
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The present sequence is that of a novel fusion protein comprising an scFv CC fragment of anti-CD7 monoclonal antibody TH-69 and domains II and III of CP Seudomonas exotoxin A (ETA), joined via a peptide linker, and including CC an N-tërminal 6xHis tag and C-terminal KDEL sequence. This novel fusion protein binds to CD7-positive T-lymphoid cells and kills them by the CC induction of apoptosis. It is an example of anti-CD7 scFv immunotoxin CC fusion proteins of the invention. An expression vector, host cells, a CC method for producing the fusion protein, and a vaccine comprising the fusion protein is used in the treatment of disorders involved a hyperproliferation of CD7-positive cells, a CC treatment or prophylaxis of Graft-versus-host disease. A claimed method CC treatment or prophylaxis of Graft-versus-host disease. A claimed method CC conditioning an animal (e.g. a human) to be transplanted with donor CC cells, tissue or organ comprises: (a) depleting the CD7-bearing cell population in (a) and/or the transplant CC donor, where the CD7-bearing cell population in (a) and/or the transplant CC protein or therapeutic composition comprising the fusion protein; and (c) introducing the transplant of the fusion or therapeutic composition comprising the fusion protein; and (c)
    Matches
                                              Query Match
                                                                                           Sequence 650
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                                                                                                                                  introducing the
                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9; Page
  98;
                         Similarity
    Conservative
                                                                                           ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; 55pp; English.
                                                                                                                                  transplant into the animal
                     90.2%;
                     Score 516; DB 6;
Pred. No. 7.1e-34;
  Mismatches
                                       Length 650;
    Indels
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Gaps
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RESULT 15
                                                                                                                    Synthetic. Chimeric.
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Region
                    11-DEC-2002; 2002WO-EP014064
                                                                    Region
                                                                                 Region
                                                                                                                                                             CD7; antibody; TH-69; scFv; exotoxin A; immunotoxin; cytostatic;
                                                                                                                                                                           Anti-CD7 antibody TH-69 scFv-Pseudomonas exotoxin A fusion protein
                                                                                                                                                                                          06-NOV-2003
                                                                                                                                                                                                                     ABR62590 standard;
     14-DEC-2001; 2001US-0339422P
                                                                                                                                  Pseudomonas
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                                                                                                                                                                                                        ABR62590;
                                               WO2003051926-A2
                                                                                                                                                       immunosuppressive;
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                                                                                                                                                                                         (first entry)
                                                             /label=
286. .64
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                                                                                                                                                       vaccine
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                                                                                  . 285
                                                                    l= Linker
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                                                              ETA
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(UYER-) UNIV ERLANGEN-NUERNBERG

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Search completed: December 23, 2004, 18:57:41 Job time: 30.8913 secs
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                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New fusion protein comprising functionally linked components of an anti-
CD7 antibody or its fragment and an immunotoxin, useful for treating CD7
positive acute T-cell and/or myeloid leukemias, or Graft-versus-host
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N-PSDB; ACF05481.
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                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                            77
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                                                                                                                      RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
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GenCore version 5.1.6
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Minimum DB
Maximum DB
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Perfect score:
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ore greater than or equal to the score of the result being printed,
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4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

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Similarity 8; Conserva DIOMTOTASSL DIOMTOTASSL	Sequence 10, Application US Patent No. 6437098 GENERAL INFORMATION: APPLICANT: SHITARA, KENYA APPLICANT: HANAI, NOBUO APPLICANT: HANGIN, NOBUO APPLICANT: MIYAJI, HIROMA APPLICANT: KUWANA, YOSHIH TITLE OF INVENTION: PROCESS FILE REFERENCE: 249-101 CURRENT APPLICATION UNMBER: FILE REFERENCE: 1995-03 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1995-05 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1995-05 PRIOR APPLICATION NUMBER: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1992-09 PRIOR FILING DATE: 1992-09 PRIOR FILING DATE: 1992-09 PRIOR FILING DATE: 1991-09 PRIOR FILING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR FILING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR FILING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19 SOFTWARE: PRIOR TILLING DATE: 1991-09 NUMBER: PRIOR TILLING DATE: 1991-09 NUMBER OF SEQ ID NOS: 19	10	8 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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100.0%; Score 572; DB 4; Length 128; iimilarity 100.0%; Pred. No. 3.5e-51; Indels 0; Gaps; Conservative 0; Mismatches 0; Indels 0; Gaps DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS	S/09225322B DRU ASA ASA SI FOR PRODUCING HUMANIZED SS FOR PRODUCING HUMANIZED US 08/454,680 5-31 US 08/408,133 3-21 US 08/292,178 US 08/292,178 9-17 JP 3-238375 9-18 JP 3-238375 9-18 JP 3-238375 9-18 JP 3-238375 9-18	ALIGNMENTS	US-08-146-206C-16 US-09-705-686-16 US-09-705-392A-16 PCT-US93-07832-16 US-08-137-117D-37 US-08-137-117D-29 US-08-436-717-29 US-08-425-763-1 US-07-934-373C-24 US-08-811-757-1 US-08-811-757-1 US-09-249-230-1 US-09-249-230-1 US-09-705-392A-24 PCT-US93-07832-24 US-08-403-853-18
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RESULT 2 US-09-225-322B-19 ; Sequence 19, Application US/09225322B

APPLICANT: SHITARA, APPLICANT: HANAI,

INFORMATION:

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APPLICANT: HANAI, NOBUO

APPLICANT: HANAI, NOBUO

APPLICANT: HASEGAWA, MAMORU

APPLICANT: HYAJI, HIROMASA

APPLICANT: HYAJI, HIROMASA

APPLICANT: KUWANA, YOSHIHISA

TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY

FILE REFERENCE: 249-101

CURRENT APPLICATION NUMBER: US/09/764,304

CURRENT FILING DATE: 2001-01-19

EARLIER APPLICATION NUMBER: US 08/454,680

EARLIER FILING DATE: 1995-03-21

EARLIER APPLICATION NUMBER: US 08/408,133

EARLIER APPLICATION NUMBER: US 08/408,133

EARLIER APPLICATION NUMBER: US 08/292,178

EARLIER APPLICATION NUMBER: US 08/292,178

EARLIER FILING DATE: 1995-03-21

EARLIER FILING DATE: 1994-08-17

EARLIER FILING DATE: 1994-09-17

EARLIER APPLICATION NUMBER: US 08/375

EARLIER APPLICATION NUMBER: US 08/375

EARLIER FILING DATE: 1994-09-18

NUMBER: OF SEQ ID NOS: 19

COPTIANTE: 1991-09-18
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APPLICANT: HASEGAWA, MAMORU
APPLICANT: HIYAII, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILING DATE: 1995-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 08/292,179
PRIOR PILING DATE: 1992-09-119
PRIOR FILING DATE: 1991-09-18
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LENGTH: 128
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NUMBER OF SEQ ID NOS: 1:
SOFTWARE: PatentIn Ver.
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APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HASSGAWA, MAMORU
APPLICANT: HASSGAWA, MAMORU
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-9
EARLIER FILING DATE: 1995-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1991-09-18
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Best Local Similarity
Matches 108; Conserv
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TYPE: PRT
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ORGANISM: Artificial Sequence
FEATURE:
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                             RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                                        DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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  RESGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                          DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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                                                                                                                                                                                       100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                       0;
                                                                                                                                                                                     Score 572; DB 4;
Pred. No. 3.5e-51;
0; Mismatches 0;
                                                                                                                                                                                                                                      Length 128;
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                                                                                                                                                                                          Indels
  128
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RESULT 5 US-09-065-059-3

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                                                                                                                                                                                                                                                                                                          RESULT 6
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                                                                                                                                                                                                                                                  Sequence 35,
Patent No. 5
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Best Local Similarity 89.8%;
Matches 97; Conservative
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                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: LIN, AUGUSTINE YEE-THARN
TITLE OF INVENTION: HUMANIZED ANTIBODIES AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 3:
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MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Bucca Ph.D., Daniel
REGISTRATION NUMBER: P-42,368
REFERENCE/DOCKET NUMBER: 50356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: OKUMURA, KO
APPLICANT: NAKATA, Motomi
TITLE OF INVENTION: THERA
COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch diskette, 1.4 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: KAYAGAKI, NO. APPLICANT: YAGITA, Hideo
                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Alexandria
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 108 amino acids
TYPE: amino acid
STRANDEDNESS: Bingle
                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
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                                                      COUNTRY:
                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                        5, Application US/08652558
5861155
                                                                                         BOSTON
                                                                      MASSACHUSETTS
                                                                                                      B: BANNER & WITCOFF
75 STATE STREET, 23RD FLOOR
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99 Canal Center Plaza
                                                        USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 96; Conserv
TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J.
TITLE OF INVENTION: MONOCLONA
TITLE OF INVENTION: CENTRAL N
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPAILIBLE
OPERATING SYSTEM: PC-DOS/MS
SOFTWARE: WordDerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/6
FILING DATE: JUNE 6, 1996
PRIOR APPLICATION DATA:
                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricla
REGISTRATION NUMBER: 27,227
REFERENCE/DOCKET NUMBER: MM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                  COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/236,520
FILING DATE: 29-APR-1994
CLASSIFICATION: 424
CLASSIFICATION: 424
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MEDIUM TYPE: Floppy disk
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STRANDEDNESS: single
TOPOLOGY: linear
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                                                             TELEPHONE:
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SYSTEM: PC-DOS/MS-DOS
                                                                 617-861-6240
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Pred. No. 3.2e-44;
5; Mismatches 6
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; MOLECULE TYPE: protein
PCT-US95-05262-2
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PCT-US95-05262-2
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Best Local S
Matches 96
                                                                                                                                        Query Match
Best Local Similarity
Matches 96; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application:
                                                                                                                                                                                                                                                                                                                     TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S.
FILING DATE: April 29, 1:
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                       NAME: Granahan, Patricia
REGISTRATION NUMBER: 27,227
REFERENCE/DOCKET NUMBER: MMN
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mayo Foundation for Medical Education Research TITLE OF INVENTION: MONOCLONAL ANTIBODIES WHICH PROMOTE TITLE OF INVENTION: CENTRAL NERVOUS SYSTEM REMYELINATION
                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 131 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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TYPE: a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US95/05262 FILING DATE:
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 81
                    61 RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 DIQMTQTTSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
                                                                                   1 DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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amino acid
                                                                  DIQMTQTTSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
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3Y: linear
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RFSGSGSGTDYSLTISNLEQEDIATYFCQQGNTLPWTFGGGTKLEIKR 128
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                                                                                                                                          Conservative
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                                                                                                                                                       87.8%;
88.9%;
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88.9%;
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29, 1994
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                                                                                                                                      Score 502; DB 5; Length 131; Pred. No. 5.2e-44; Indels 3; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 502; DB 1; Length 131; Pred. No. 5.2e-44; 3; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Smithy & Reynolds, P.C
                                                                                                                                      0;
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US-09-386-658A-4

Sequence 4, Application US/09386658A
Patent No. 6593137
GENERAL INFORMATION:
APPLICANT: Erlanger, Bernard F.
APPLICANT: Chen, Bi-Xing
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR FULLERENES
FILE REFERENCE: 0575/54182

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US-08-487-761-13
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                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-487-761-13
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Patent No. 62178
                                                                                                                                    Matches
                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/487,761
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/086,411
FILING DATE: 29-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 TELEFAX: (215) 454-380
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cheadle, Christoph
APPLICANT: South, Victoria J.
TITLE OF INVENTION: Monoclona
TITLE OF INVENTION: Epidermal
TITLE OF INVENTION: Employing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acid
                                                                                                                                                                                                                                                                                                                                NAME: GOOdman, ROBANNE
REGISTRATION NUMBER: 32,534
REFERENCE/DOCKET NUMBER: A0:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 454-3817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: Macintosh
OPERATING SYSTEM: Syste
SOFTWARE: Word 5.0 (Pat
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Rhone-Poulenc Rorer Inc
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                          61
61 RFSGSGSGTDYSLTISNLEPEDIATYYCQQYSKIPYTFTGGTKLEIKR 108
                                                                                                                                    94;
                                                                                 Similarity
                   RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                                                  EIHMTQTTSSLSASLGDRVTISCSASQDIRNYLNWYQQKPDGTVKLLIYYTSTLHSGVPS
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                                                                                                                                                                                                                                                                 112 amino acids
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Kris, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Givol, David
                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                    (215) 454-3808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ricca, George A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schlessinger, Joseph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                87.1%;
87.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal Antibodies Specific to Human
Epidermal Growth Factor Receptor and Therapeutic Methods
Employing Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        System 7.1
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                                                                                                                                Score 498; DB 3;
Pred. No. 1.1e-43;
6; Mismatches 8
                                                                                                                                                                 DB 3; Length 112;
                                                                                                                                    Indels
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                                                                                                                                    Gaps
                                                                   60
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; LENGTH: 109
; TYPE: PRT
; ORGANISM: MOUSE
US-09-386-658A-4
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                                                                                                                                ; MOLECULE TYPE: protein US-08-458-516-9
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                                                          Matches
                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 4
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                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/059,159

FILING DATE: 03-MAY-1993

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 11823-37-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/386,658A CURRENT FILING DATE: 1999-08-31 NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                           TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Co, Man Sung
APPLICANT: TBO, J. Yun
TITLE OF INVENTION: Humanized F
                                                                                                                                                                LENGTH: 107 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: One Market I
                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 RFSGSGSGTDYSLTINNLEPEDIATYFCQQYSRLPFTFGSGTKLEIKR 108
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                                                        l Similarity
93; Conserv
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DIQMTQTASSLFASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS 60
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                                                                                                                                                                                                                                                                           415-326-2400
                                                                       85.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/08/458,516
                                                                                                                                                                                                                                         9
                                                                                                                                                                                                                                                                                                               11823-37-3
                                                    Score 488; DB 1; Length 107; Pred. No. 1.1e-42; Mismatches 10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 497; DB 4;
Pred. No. 1.3e-43;
8; Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Version #1.25
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                                                        Gaps
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US-08-649-100-17

Patent No.

Sequence 17, Application US/08649100 Patent No. 6114507

GENERAL INFORMATION:
APPLICANT: SHIRAKA
APPLICANT: MATUSUE
APPLICANT: NAGATA,

SHIRAKAWA, KAMON MATUSUE, TOMOKAZU NAGATA, SHIGEKAZU

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RESULT 13
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US-08-458-516-5
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APPLICANT: Co, Man Sung
APPLICANT: T80, J. Yun
TITLE OF INVENTION: Human
TITLE OF INVENTION: GPIIE
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5,
Patent No. !
                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 127 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 03-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30.22
                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                        TOPOLOGY: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
                                                                                                                                                                                                                  local Similarity
                                                                       61 RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIK 107
                                                                                                                           21 DIOMTOTTSSLSASLGDRVTISCRASQDINNYLNWYQQKPDGIVKLLIYYTSTLHSGVPS
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                                                                                                                                             1 DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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                                                     RESGSGSGTDYSLTISNLEQEDIATYFCQQGNTLPWTFGGGTKLEIK 127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08458516
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                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                        protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                                                                                  85.3%;
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                                                                                                                                                                                             Score 488; DB 1;
Pred. No. 1.3e-42;
4; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                          11823-37-3
                                                                                                                                                                                                  10; Indels
                                                                                                                                                                                                                                   Length 127;
                                                                                                                                                                                                 0
                                                                                                                                                                                                 Gaps
                                                                                                                            80
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APPLICANT: VASQUEZ, MAXIMILIANO
TITLE OF INVENTION: ANTI-FAS LIGAND ANTIBODY AND ASSAY
TITLE OF INVENTION: METHOD USING THE ANTI-FAS LIGAND ANTIBODY

NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:

41

STREET: PO BOX 747 CITY: FALLS CHURCH

E: BIRCH, STEWART, KOLASCH AND BIRCH
PO BOX 747

ADDRESSEE:

```
Sequence 7, Application US/09199149

Fatent No. 6150099

GENERAL INFORMATION:

APPLICANT: Jonak, Zdenka L.

APPLICANT: Taylor, Alexander H.

APPLICANT: Trulli Jr., Stephen H.

APPLICANT: Johanson, Kyung O.

ITILE OF INVENTION: Humanized Monoclonal Antibodies

FILE REFERENCE: P50860

CURRENT APPLICATION NUMBER: US/09/199,149

CURRENT FILING DATE: 1998-11-24

NUMBER OF SEQ ID NOS: 37

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 7

LENGTH: 108
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                                                                             ; TYPE: PRT ; ORGANISM: murine cells US-09-199-149-7
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Best Local S
Matches 94
Query Match
Best Local Similarity
Matches 92; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/649
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MURPHY JR, GERALD M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1110
TELECOMMUNICATION INFORMATION:
TELECHONE: (703) 205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELBEAX: (703) 205-8050
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 127 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
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 Score 487; DB 3; Length 108; Pred. No. 1.4e-42; 6; Mismatches 10; Indels
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US-09-813-659-30
Search completed: December 23, Job time: 8.13276 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 30436.18USD2
CURRENT APPLICATION NUMBER: US/09/813,659
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 09/549,067
PRIOR FILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: 08/539,436
PRIOR APPLICATION NUMBER: 08/539,436
PRIOR FILING DATE: 1993-09-13
PRIOR FILING DATE: 1993-09-13
PRIOR FILING DATE: 1993-09-13
                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver.
SEQ ID NO 30
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bajorath, Jurgen
APPLICANT: Fell, H. Perry
APPLICANT: Gilliland, Lisa K.
APPLICANT: Gilliland, Lisa K.
TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
TITLE OF INVENTION: FUSION PROTEINS IN A MAMMALIAN CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 08/013,420 PRIOR FILING DATE: 1993-02-01
                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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                                                                           RFSGSGSGTDYSLTIANLQPEDIATYFCQQGNTLPWTFGGGTKLVTKR
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Linsley, Peter S.
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                 2004, 19:08:02
                                                                                                                                                                                                                            Score 487; DB 4; Length 274; Pred. No. 4.3e-42; Mismatches 11; Indels
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB seq length: 200000000
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Published Applications AA:*

1: /cgn2_6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

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7: /cgn2_6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

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11: /cgn2_6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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491	494	497	502	502	510	510	572	572	572	572	572	572	Score
85.8	86.4	86.9	87.8	87.8	89.2	89.2	100.0	100.0	100.0	100.0	100.0	100.0	Query
107	107	109	131	131	108	108	128	128	128	128	128	128	Query Match Length
17	17	14	14	œ	14	9	14	14	14	14	9	9	8
US-10-473-977-69	US-10-741-657A-16	US-10-197-080-4	US-10-010-729-64	US-08-779-784-21	US-10-234-671-10	US-09-056-160B-10	US-10-166-626-19	US-10-166-626-10	US-10-265-713-19	US-10-265-713-10	US-09-764-304-19	US-09-764-304-10	ID,
Sequence 69, Appl	Sequence 16, Appl	Sequence 4, Appli		Sequence 21, Appl	Sequence 10, Appl	Sequence 10, Appl	Sequence 19, Appl	Sequence 10, Appl	Sequence 19, Appl	e 10,	Sequence 19, Appl	Sequence 10, Appl	Description

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14
484	484	484	485	485	487	487	487	487	487	487	487	487	487	488	488	488	488	488	488	488	488	488	488	488	488	488	488	488	488	488	488
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127	107	107	107	107	555	504	302	302	302	302	274	274	108	657	656	643	642	638	637	601	127	108	107	107	107	107	107	107	107	107	107
14	14	14	15	14	15	14	14	14	9	9	14	9	14	16	16	16	16	16	16	9	14	13	16	16	16	16	16	15	15	15	15
US-10-268-883-5	-1	-10-269-010-	-10	US-10-310-674A-34	-10-107-991E	US-10-207-655-348		L	US-09-813-659-32	US-09-813-659-18	US-10-283-610A-30	US-09-813-659-30	1	US-10-296-085A-68	w	-10-296	-10-296-085A-3	10-296-085A-		0	-10-084	-10-140-555-	-10-410-913-5	v	-10-411-012-	-10-410-997-	-10-410-930-5	-10-411-049-5	US-10-410-962-52	-10-411-026-	-10-411-037-5
رن ر	е б,	Sequence 2, Appli	e 10,	е 34,	e 3,	346	32	e 18,	Sequence 32, Appl	18,	.0	30,	7,	89	Sequence 39, Appl	69,	38,	21,	e 1		N	4	s	u	e 52,	52,	52,	Sequence 52, Appl	e 52,	-	52,

ALIGNMENTS

; FEALUKE: ; OTHER INFORMATION: CDNA KM-641 US-09-764-304-10	; ORGANISM: Artificial Sequence	TYPE: PRT	LENGTH: 128	SECTION 10		; EARLIER FILING DATE: 191-09-18		; EARLIER FILING DATE: 1992-09-17	; EARLIER APPLICATION NUMBER: US07/947,674	FILING DATE: 1994-08-17	APPLICATION N	FILING DATE:	APPLICATION	FILING DATE: 1995-05-3	APPLICATION NUMBER: US	FILING DATE	APPLICATION		CURRENT APPLICATION NUMBER: US/09/764,304	; FILE REFERENCE: 249-101	z		••	••	: HANAI, N		; GENERAL INFORMATION:	; Patent No. US20020026036A1	; Sequence 10, Application US/09764304	US-09-764-304-10	RESITT	HUMANIZED CHIMERA ANTIBODY	RESULT 1 US-09-764-304-10 ¡Sequence 10, Application US/09764304 Patent No. US20020026036A1 GENERAL INFORMATION: APPLICANT: HANAI, NOBUO APPLICANT: HASEGAWA, MAMORU APPLICANT: KUWANA, VOSHHIISA APPLICANT: KUWANA, VOSHHIISA APPLICANT: KUWANA, VOSHHIISA CURRENT FILING DATE: 249-101 CURRENT FILING DATE: 2001-01-19 EARLIER APPLICATION NUMBER: US/09/764,304 CURRENT FILING DATE: 1995-05-31 EARLIER APPLICATION NUMBER: US 08/454,680 EARLIER APPLICATION NUMBER: US 08/454,680 EARLIER APPLICATION NUMBER: US 08/408,133 EARLIER APPLICATION NUMBER: US 08/225,322 EARLIER APPLICATION NUMBER: US 08/292,178 EARLIER FILING DATE: 1995-03-21 EARLIER FILING DATE: 1992-09-17 EARLIER APPLICATION NUMBER: US 08/292,178 EARLIER FILING DATE: 1992-09-17 EARLIER FILING DATE: 1992-09-17 EARLIER FILING DATE: 1991-09-18 NUMBER OF SEQ ID NOS: 19 SEQ ID NO 10 LENGTH: 128 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: CDNA KM-641 US-09-764-304-10	
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Sequence 10, Application US/10265713
Publication No. US20030095964A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAAI, HIROMASA
APPLICANT: MIYAAI, HIROMASA
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CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1994-08-17
EARLIER FILING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US 08/292,774
EARLIER FILING DATE: 1994-08-17
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Best Local Similarity
Matches 108; Conserv
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SEQ ID NO 19
LENGTH: 128
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Best Local Similarity
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APPLICANT: KUMANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
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NUMBER OF SEQ ID NOS: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: light chain OTHER INFORMATION: variable region
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ORGANISM: Artificial Sequence
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KUWANA, YOSHIHISA
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APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR FILING DATE: 1999-01-05
PRIOR FILING DATE: 1999-01-05
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/202,178
PRIOR PILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 08/292,178
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SEQ ID NO 10
LENGTH: 128
TYPB: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-10-265-713-10
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CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT APPLICATION NUMBER: US/09/225,322
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR PPLICATION NUMBER: US 08/454,680
PRIOR PLICATION NUMBER: US 08/454,680
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US07/947,674
                                                         SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 19
LENGTH: 128
TYPE: PRT
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Best Local
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APPLICANT: HANAI, I
APPLICANT: HASEGAWI
                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: JP 3-238375 PRIOR FILING DATE: 1991-09-18
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ORGANISM: Artificial Sequence FEATURE:
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HASEGAWA, MAN
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OTHER INFORMATION: Description of Artificial Sequence:light chain

7.12

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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/166,626
CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR FILING DATE: 1999-01-05
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
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                                                                      Sequence 19, Application US/10166626 Publication No. US20030166876A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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SEQ ID NO 10
LENGTH: 128
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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAM
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
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; OTHER INFORMATION: Description of Artificial Sequence:light chain ; OTHER INFORMATION: variable region US-10-166-626-19
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US-09-056-160B-10
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PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR PILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR PILING DATE: 1992-09-17
PRIOR FILING DATE: 1992-09-17
PRIOR FILING DATE: 1991-09-18
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Patent No. U
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APPLICANT: Baca, Manuel
APPLICANT: Wells, James A.
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TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lowman, Henry B.

APPLICANT: Chen, Yvonne M.

TITLE OF INVENTION: ANTI-VEGF ANTIBODIES

NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 128
                                          APPLICATION NUMBER: US/09/056,160B
FILING DATE: 06-Apr-1998
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech
APPLICATION NUMBER: FILING DATE: 06-AUG
                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 DIQMTQTASSLPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/09056160B
                                                                                                                                                                                                                                                                                                                                                                                     1 DNA Way
                                                                                                                                                                                                                                                                                                                USA
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UMBER: 60/054,856
06-AUG-1997
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ATTORNEY/AGENT INFORMATION:

P1093R2

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US-10-234-671-10
, Sequence 10, Application US/10234671
; Publication No. US20030190317A1
; Publication No. US20030190317A1
; GENERAL INFORMATION:
; APPLICANT: Baca, Manuel
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REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P109
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acids
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Best Local Similarity
                                          INFORMATION FOR SEQ ID NO: 10:
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TOPOLOGY: Lin-
056-16^
                                                                                           APPLICATION NUMBER: 09/056160

PILING DATE: 06-APR-1998

APPLICATION NUMBER: 60/126446

FILING DATE: 07-APR-1997

APPLICATION NUMBER: 60/054856

FILING DATE: 06-AUG-1997

ATTORNEY/AGENT INFORMATION:

NAME: Cui, Steven X.

REGISTRATION NUMBER: 44,637

REFERENCE/DOCKET NUMBER: P1093R2C1

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61
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STATE: California
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                                                           TELEPHONE: 650/225-8674
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1 DNA Way
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LENGTH:
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amino acids
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Pred. No. 1.6e-39;
7; Mismatches 6
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US-08-779-784-21
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                                                                                                                      TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-779-784-21
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Publication No. US20020164325A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 89.2%;
Best Local Similarity 88.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                               Best Local
Matches 9
                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/779,784
FILING DATE: 07-JR4.1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/692,084
FILING DATE: 08-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/236,520
FILING DATE: 08-AUG-1996
PRIOR APPLICATION NUMBER: US 08/236,520
FILING DATE: 29-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                 Query Match
                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rodriguez, Mos
APPLICANT: Miller, David
APPLICANT: Asakura, Kunik
                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: PROMOTION OF CENTRAL NERVOUS SYSTEM
TITLE OF INVENTION: REMYELINATION USING MONOCLONAL AUTOANTIBODIES
                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                              TELEPHONE: 201-343-1684
                                                                                                                                                                                                                                                                                                        NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1199-1-001 CIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 411 Hacke
STREET: Floor
CITY: Hackensack
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ZIP: 0760:
                                                                                                                                                                                          LENGTH:
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1 DIQMTQTASSLFASLGDRVTISCSASQDISNYLNWYQQKFDGTVKLLIFYSSNLHSGVFS
                                                 96;
                                                                Similarity
                                                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                              131 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                 Conservative
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                                                                                                                                                                                                                                                                       201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                              87.8%;
                                               ω,
                                               Score 502; DB 8;
Pred. No. 1e-38;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 510; DB 14;
Pred. No. 1.6e-39;
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Continental Plaza, 4th
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                                                                                    8
                                                  9
                                                                                 Length 131;
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                                                  Indels
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; LENGTH: 109
; TYPE: PRT
; ORGANISM: Mouse
US-10-197-080-4
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                                                                                   APPLICANT: Erlanger, Bernard F.
APPLICANT: Sheetz, Michael
APPLICANT: Sheetz, Michael
APPLICANT: Brus, Louis
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NANOTUBES AND RELATED METHODS AND COMPOSI
FILE REFERENCE: 0575/67096-A
CURRENT APPLICATION NUMBER: US/10/197,080
CURRENT FILING DATE: 2002-07-16
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/10197080 Publication No. US20030113940A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 64, Application US/10010729
Publication No. US20030185827A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Miller, David J.
APPLICANT: Pease, Larry R.
APPLICONT: Pease, Larry R.
TITLE OF INVENTION: Human IgM Antibodies and Diagnostic and
TITLE OF INVENTION: Therapeutic Uses Thereof Particularly in the Central Nervous
TITLE OF INVENTION: System
FILE REFERENCE: 1199-1-005CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1996-08-08
PRIOR APPLICATION NUMBER: 08/236,520
PRIOR FILING DATE: 1994-04-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/010,729
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR FILING DATE: 2000-12-05
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/322,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 08/692,084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1999-05-28 PRIOR APPLICATION NUMBER: 08/PRIOR FILING DATE: 1997-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J
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Pred. No. 1e-38;
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US-10-473-977-69
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US-10-473-977-69
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US-10-741-657A-16
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                Query Match 85.8%;
Best Local Similarity 87.9%;
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Best Local
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TITLE OF INVENTION: ANTIBODIES AGAINST GPR64 AND USES THEREOF
FILE REFERENCE: 05882.0177.NPUS01
CURRENT APPLICATION NUMBER: US/10/741,657A
CURRENT FILING DATE: 2003-12-19
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin version 3.2
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NUMBER OF SEQ ID NOS: 72
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TYPE: PRT
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TYPE: PRT
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No. US20040197325A1
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Conservative
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                Score 491; DB 17;
Pred. No. 8.8e-38;
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Pred. No. 4.6e-38;
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Pred. No. 2.5e-38;
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PRIOR FILING DATE: 2001-10-10
PRIOR PELICATION NUMBER: US 60/344,692
PRIOR FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: US 60/387,292
PRIOR FILING DATE: 2002-06-07
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR APPLICATION NUMBER: US 60/396,594
PRIOR PILING DATE: 2002-06-25
PRIOR PILING DATE: 2002-07-17
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Best Local (
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SEQ ID NO 52
                                                                                                                 GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
APPLICANT:
                                                      APPLICANT: Neose Technologies, Inc. APPLICANT: DeFrees, Shawn APPLICANT: Zopf, David
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/404,249 PRIOR FILING DATE: 2002-08-16
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APPLICANT: DeFrees, Shawn
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NUMBER OF SEQ ID NOS: 75
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Bowe, Caryn
              Bayer, Robert
Hakes, David
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Pred. No. 1.7e-37;
4; Mismatches 10;
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PRIOR FILLING DATE: 2001-10-10
PRIOR PELLING DATE: 2001-10-10
PRIOR PELLING DATE: 2001-10-19
PRIOR PELLING DATE: 2001-10-19
PRIOR PELLING DATE: 2002-06-07
PRIOR PELLING DATE: 2002-06-07
PRIOR PELLING DATE: 2002-06-25
PRIOR PELLING DATE: 2002-07-17
PRIOR PELLING DATE: 2002-07-17
PRIOR PELLING DATE: 2002-07-17
PRIOR PELLING DATE: 2002-08-16
PRIOR PELLING DATE: 2002-08-28
PRIOR PELLING DATE: 2002-08-28
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CURRENT APPLICATION NUMBER: US/10/411,026
CURRENT FILING DATE: 2003-04-09
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                                                                                                                                                                                                                                                                                                                          LENGTH:
  61
                        61 RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIK 107
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                                                                                      1 DIQMTQTTSSLSASLGDRVTISCRASQDINNYLNWYQQKPDGIVKLLIYYTSTLHSGVPS
                                                                                                                                                                                             Similarity
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RESGSGSGTDYSLTISNLEQEDIATYFCQQGNTLPWTFGGGTKLEIK 107
                                                                                                                                                                           Conservative
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                                                                                                                                                                       Score 488; DB 15;
Pred. No. 1.7e-37;
4; Mismatches 10;
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Maximum Match 100%
Listing first 45 summaries
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Copyright (c) 1993 - 2004 Compugen Ltd.
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A;Residues: 1-10
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A;Note: sequence extracted from NCBI backbone (NCBIP:60876)
R;Puccetti, A.; Koizumi, T.; Migliorini, P.; Andre-Schwartz
J. Exp. Med. 171, 1919-1930, 1990
A;Title: An immunoglobulin light chain from a lupus-prone muhare: PLO220, MUID:90278348; PMID:1693654
A;Recession: PLO220
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Molecule type: mRNA
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J. Biol. Chem. 266, 6607-6613, 191
J. Biol. Chem. 266, 6607-6613, 191
A;Title: Heavy and light chain variable region sequences and antibody properties of a A;Reference number: A38740; MUID:91177923; PMID:1706720
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C;Superfamily: immunoglobulin V region; immun
C;Keywords: heterotetramer; immunoglobulin
E;19-93/Domain: immunoglobulin homology <IMM>
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A; Title: Molecular heterogeneity of auto-anti-idiotypic
A; Reference number: A49026; MUID:91364791; PMID:1909645
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Ig kappa chain V region (Py69) - mouse (Species: Mus musculus (house mouse) (Species: Mus musculus (house mouse) (C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 09-Jul-2004 C;Accession: G38740 R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R. J. Biol. Chem. 266, 6607-6613, 1991 A;Title: Heavy and light chain variable region sequences and antibody prope A;Reference number: A38740; MUID:91177923; PMID:1706720 A;Accession: G38740
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A;Title: B-cell proliferation initiated by Ia cross-linkin A;Reference number: S69900; MUID:92165291; PMID:1537587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig kappa chain (clone KL2.18 / KL4B10 / C;Species: Mus musculus (house mouse) C;Date: 14-Feb-197 #sequence revision C;Accession: S69900; S69907; S69908
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A;Molecule type: DNA
A;Residues: 1-108 <WY3>
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A;Cross-references: UNIPROT:Q91WS9
C;Superfamily: immunoglobulin V region;
                                                             A; Molecule type: mRNA
                                                                                 A; Status: preliminary; nucleic acid
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A; Residues: 1-108 < WYS>
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88.9%;
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Pred. No. 1.6e-39;
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Pred. No. 4.7e-39;
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PMID:1537587
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Ig kappa chain precursor V region (AC-1001) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 21-Jan-2000
C;Accession: A29380
R;Chen, H.T.; Kabat, E.A.; Lundblad, A.; Ratcliffe, R.M.
J. Biol. Chem. 262, 13579-13583, 1987
A;Title: Nucleotide and translated amino acid sequences of cDNA coding for the code of and translated amino acid sequences of cDNA coding for the code of the code of the code of the code of cDNA coding for the code of cDNA code of cDNA coding for the code of cDNA 
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A;Cross-references: UNIPROT:P01643
C;Comment: This chain was isolated from a myeloma protein.
C;Comment: An immunoglobulin heterotetramer subunit consists of two identical light hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate int C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer
F;16-90/Domain: immunoglobulin homology <IMM>
F;23-88/Disulfide bonds: #status predicted
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Bur. J. Biochem. 59, 525-537, 1975

Bur. J. Biochem. 59, 525-537, 1975

A;Title: Determination of the primary structure of a mouse IgG2a immunoglobulin.

A;Reference number: A01926; MUID:76091934; PMID:812696

A;Accession: A01926

A;Accession: A01926
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A29380
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R; Schiff, C.; Fouge:
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;Species: Mus musculus (house mouse)
;Date: 24-Apr-1984 #sequence_revision
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Pred. No. 4.9e-39;
7; Mismatches 7
  Score 495; DB 2;
Pred. No. 2.4e-38;
3; Mismatches 10
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Pred. No. 1.1e-38;
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C;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #rear_...
C;Accession: JL0080
R;Kaartinen, M.; Rocca-Serra, J.; Maekelae, O.
Mol. Immunol. 25, 859-865, 1988
A;Title: Combinatorial association of V genes: one VH gene
A;Reference number: JL0076; MUID:89096973; PMID:3211160
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S69903
A;Cross references: GB:M27793; NID:g197161; PIDN:AAA38937.1; PI A;Note: the authors translated the codon AGG for residue 30 as C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: immunoglobulin F;1-6/Domain: signal sequence (fragment) #status predicted <SIG F;7-115/Product: Ig light chain #status predicted <MAT> F;22-96/Domain: immunoglobulin homology <IMM> F;20-40/Region: complementarity-determining 1 F;56-62/Region: complementarity-determining 2
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A;Title: B-cell proliferation initiated by Ia cross-li A;Reference number: S69900; MUID:92165291; PMID:153758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Date: 14-Feb-1997 #sequence_revision 13-Mar-1997 #text_change
C;Accession: S69903; S69904; S69905
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                                                                                                                                                                                                                                                                                                                                                                                        Ig kappa chain precursor V region (anti-phenyloxazolone, 18C10) C;Species: Mus musculus (house mouse) C;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: EMBL:X55046; NID:g511033; PIDN:CAA38886.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology F;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary; translation not shown A;Molecule type: DNA A;Residues: 1-108 <W3>
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A; Residues: 1-108 < WY2>
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A;Status: preliminary; translation
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A; Residues: 1-115 < KAA>
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Pred. No. 3.1e-38;
4; Mismatches 9
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PMID:1537587
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Arg, 84-Ser, and 93-Ala
C;Comment: The sequence shown is HP R16.7.
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer
F;16-90/Domain: immunoglobulin homology <IMM>
F;23-88/Disulfide bonds: #status pradicted
F;23-88/Disulfide bonds: #status pradicted
                                                                                                                                                                                                           C;Species: Mus musculus (house mouse)
C;Date: 09-Oct-1992 #sequence_revision(
C;Accession: PHO888
R;Shalaby, M.R.; Shepard, H.M.; Presta,
J. Exp. Med. 175, 217-225, 1992
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C;Date: 06-Jul-1982 #sequence_revision 06-Jul-1982 #text_change 09-Jul
C;Accession: A01927
R;Siegelman, M.; Capra, J.D.
Proc. Natl. Acad. Sci. U.S.A. 78, 7679-7683, 1981
A;Title: Complete amino acid sequence of light chain variable regions A;Reference number: A01927; MUID:82150934; PMID:6801658
A;Accession: A01927
                                                                         444005
                                                                                                                                                                          A;Title: Development of humanized bispecific antibodies A;Reference number: PH0885; MUID:92113462; PMID:1346155
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A;Experimental source: strain A/J
A;Note: hp 93G7 differs in having 93-Met; HP 123E6 differs in having 7-Ser, 92-Tyr,
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Best Local S
Matches 95
                                                                                    Superfamily: immunoglobulin V region; Keywords: immunoglobulin
                                                                                                                                         Molecule type: mRNA
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   Matches
                  Best
                                                                     16-90/Domain:
                                                                                                                          Residues: 1-109 <SHA>
                                                                                                                                                           Accession: PH0888
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Best Local :
                                  Query Match
                  Local Similarity
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                                                                   immunoglobulin
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                85.3%;
86.1%;
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Pred. No. 3.4e-38;
                Score 488; DB 2;
Pred. No. 9.1e-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 489; DB 1;
Pred. No. 7.3e-38;
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Mismatches
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                                                                                                        immunoglobulin
                                                                     <IMM>
                                                                                                                                                                                                                             Rodrigues, M.L.; Beverley, P.C.L.;
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                                Length 109;
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Ig kappa chain precursor V region (5-27) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 27-Jul-1990 #sequence_revision 27-Jul-1990 #text_change 09-Ju
C;Accession: A34904
R;Bedzyk, W.D.; Herron, J.N.; Edmundson, A.B.; Voss Jr., E.W.
J. Biol. Chem. 265, 133-138, 1990
A;Title: Active site structure and antigen binding properties of idic
A;Seference number: A34904
A;Seference number: A34903; MUID:90094387; PMID:2104617
A;Accession: A34904
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-126 <BED>
A;Cross-references: UNIPROT:Q91WF8
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;35-109/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V region - mouse
C;Species: Mus musculius (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
C;Accession: S38862
R;Fischer, R.; Voss, A.; Hunziker, W.; Stierhof, Y.D.; Kreuzaler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT
S38862
                                                                                                                                                                                                                   A;Cross-references: EMBL:X75854; NID:g429109; PID:g429110 (;Superfamily: immunoglobulin V region; immunoglobulin hot C;Keywords: heterotetramer; immunoglobulin F;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-108 <FIS>
                                                                                                                                                                                                                                                                                                                                                                                R;Fischer, R.; Voss, A.; Hunziker, W.; Stierhof, Y.D.; Kreuzaler, submitted to the EMBL Data Library, August 1993
A;Description: Production and cloning of TMV-specific monoclonal
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                                                                         DVQMTQTTSSLAASLGDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS
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Pred. No. 1.5e-37;
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Pred. No. 1.1e-37;
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RESULT 15
A26406

A26406

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A26406

C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Species: Jo-Un-1989 #sequence_revision 30-Jun-1991 #text_change 09-Jul-2004
C;Accession: A26406
R;Sanz, I; Capra, J.D.
Proc. Natl. Acad. Sci. U.S.A. 84, 1085-1089, 1987
A;Title: V-K and J-K gene segments of A/J Ars-A antibodies: somatic recombination general A;Accession: A26406
A;Molecule type: DNA
A;Residues: 1-128 <ANN-A;Residues: 1-12
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,
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   RA Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., RA Richards S., Worley K.C., Hale S., Garcia A.M., Gunaratne P.H., RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., RA Miting M., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G., RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E., The Mouse C.DNA sequences.",

"Generation and initial analysis of more than 15,000 full-length human for the first of the 
                 EMBL; BC013496; AAH13496.1; --
PIR; A38740; A38740.
PIR; C38740; C38740.
PIR; B38740; G38740.
PIR; B38740; G38740.
InterPro; IPR00310; Ig-1ike.
InterPro; IPR003150; Ig_MHC.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR00356; Ig_V.
Pfam; PF007654; C1-8et; 1.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
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Q91WS9;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
Q1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
Q1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein (Fragment).
Mus musculus (Mouse)
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STRAIN=FVB/N; TISSUE=Colon;
MEDLINE=22388257; PubMed=12477932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. NCBI_TaxID=10090;
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STRAIN=FVB/N; TISSUE=Colon;
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KV1W HUMAN
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P01649 mus musculu
Q96pf6 homo sapien
Q6pih7 homo sapien
Aah34141 homo sapien
P01595 homo sapien
P01616 homo sapien
P04431 homo sapien
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P01600 homo sapien
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Result

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KV5K MOUSE
KV5L MOUSE
AAR10992
KV5N MOUSE
KV5N MOUSE
KV5M MOUSE

Q91WF8

400 400 399

11088 11088

OSUL84
KVSU MOUSE
AAR11052
AAR11043
AAR11043
AAR11040
KV1B HUMAN
KV1O HUMAN
KV1Y HUMAN
KV1A HUMAN
KV1A HUMAN
KV1A HUMAN
OSUL70
O96SA9

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Q9UL77
KV5G MOUSE
Q6GMX9
Q6GMX9
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KV1R HUMAN
Q6GMW1
KV1L HUMAN
Q7Z3Ÿ4

KV1M_HUMAN

Minimum Maximum

BG 1

geq pea

length: 0 length: 2000000000

Post-processing:

Minimum Match 0% Maximum Match 100% Listing first 45 s

summaries

Total number of hits satisfying chosen parameters:

1825181 segs, 575374646 residues

Searched:

Database

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Sequence: Title: Perfect score:

US-10-089-500-56 572 1 DIQMTQTASSLPAS

Scoring table:

BLOSUM62

Gapop 10.0 ,

Gapext 0.5

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OM protein - protein search, using

8 mode1 Copyright

GenCore version 5.1.6 (c) 1993 - 2004 Compugen

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RESULT 3
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Q8R062
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SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 1.
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21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                         InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                              PIR; A01926; KVMS73.
HSSP; P01594; 1JV5.
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Mammalia; Eutheria;
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Mus musculus (Mouse)
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                                                                                                                                                                                                                                                        Direct
                                                                                                                                                                                                                                                                                                                                         "Determination of the primary structure of immunoglobulin. Amino-acid sequence of the Eur. J. Biochem. 59:525-537(1975).
                                                                                                                                                                                                                                                                                                                                                                      Schiff C.,
                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=76091934; PubMed=812696;
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                                                                                                Similarity
                                                                                RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWIFGGGTKLEIKR
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PRELIMINARY;
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Rodentia;
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25781 MW;
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                                                                                                                     Score 498; DB
Pred. No. 6.6e
9; Mismatches
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Pred. No. 2
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By similarit
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Complementarity-determining-3.
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Complementarity-determining-2.
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                                                                                                                                                                                                                                               Framework-1
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PRT;
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                                                                                                                                                            2AD29D92A72AA0A3 CRC64;
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234
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light chain.";
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KV5K_MOUSE
ID KV5K MOUSE
AC PO1644;
DT 21-JUL-1986
DT 21-JUL-2096
DT 05-JUL-2004
DE 1g kappa chai

STANDARD;

108 B

kappa chain

(Rel. (Rel.

01, Created)
01, Last sequence update)
44, Last annotation update)
V region HP R16.7.

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                                                                                                                                           Query Match
Best Local S
Matches 94
                                                                                                                                                                                                                                                                                                        HSSP; P01594; 1JV5.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; Cl-set; 1.
Pfam; PF07654; Cl-set; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Krzywinski M.1.,
Jones S.J., Marra M.A.;
"Generation and initial analysis of
"Generation by and mouse cDNA sequences.";
"And mouse cDNA sequences.";
                                                                                                                                                                                                                 SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
Hypothetical protein.
SEQUENCE 234 AA; 25858 MW; 4EB08
                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (APR-2002) to the EMBL; BC027418; AAH27418.1; HSSP; P01594; LJV5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=FVB/N; TISSUE=Colon;
MEDLINE=22388257; PubMed=12477932;
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Mammalia; Eutheria;
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01-JUN-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM
STRAIN=FVB/N;
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81
                                   61
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                                                                                                                                                                Similarity
                                                                                          DIQMTQTASSLFASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPS
                        RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                       DIQMTQTTSSLSASLGDRVTISCSASQGISNYLNWYQQKPDGTVKLLIYYTSSLHSGVPS
RFSGSGSGTHYSLTISNLEPEDIATYYCQQYSQFPFTFGSGTKLEIKR
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i; TISSUE=Colon;
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TISSUE=Colon;
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Rodentia;
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Last annotation updat
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                                                                                                                                           Score 494; DB 2; I
Pred. No. 4.1e-42;
6; Mismatches 8;
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RESULT 5

RV5L MOUSE
ID "KV5L M
AC P01645
DT 21-JUL
DT 21-JUL
DT 05-JUL
DT 05-JUL
OT 05-Was man
OC Eukary
OC Mammal
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Best Local S
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21-JUL-1986 (Rel
21-JUL-1986 (Rel
05-JUL-2004 (Rel
15 kappa chain V
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HSSP; P01594; 1JV5.
InterPro; IPR007110;
InterPro; IPR003596;
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DISULFID
NON_TER
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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DOMAIN
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"Complete amino acid sequence of light chain variable regions derived from five monoclonal anti-p-azophenylarsonate antibodies differing with respect to a crossreactive idiotype.";
proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
-!- MISCELLANEOUS: Anti-arsonate hybridoma protein.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
HSSP; P01607; 1REI. Interpro; IPR0077110; Ig-like. Interpro; IPR003596; Ig_v. Pfam; PF00047; Ig; 1. SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                STRAIN=A/J;
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
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SMART; SM00406; IGV;
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(Rel.
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1 23 Framework-1.
24 34 Complementarity-determin
35 49 Framework-2.
50 56 Complementarity-determin
57 88 Framework-3.
89 97 Complementarity-determin
98 108 Framework-4.
23 88 By similarity.
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. 01, Last sequence 44, Last annual
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Pred. No. 5.4e-42;
3; Mismatches 10;
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Complementarity-determining-3. Framework-4.
By similarity.
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Sciurognathi; Muridae;
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Best Local &
KV5N MOUSE STANDARD; PRT; 108
P01647;
P01647;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence upda
05-JUL-2004 (Rel. 44, Last annotation up
1g kappa chain V-V region HP 124E1.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata;
Bukaryota; Metazoa; Rodentia; Sciurognat
                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kresl
"Antinuclear autoantibodies fro
Submitted (SEP-2003) to the EMF
""" AY436832; AAR10992.1; -.
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MAR-2004 (TrEMBLrel. 27, Last sequence update)
MAR-2004 (TrEMBLrel. 27, Last snotation update)
immunoglobulin kappa light chain (Fragment).
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s from B6.Sle1 mice.
e EMBL/GenBank/DDBJ
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Pred. No. 6.9e-42;
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 Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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InterPro; IPRO03596; Ig v.
Pfam; PPRO047; ig; 1.
SMART; SM00406; IGv; 1.
Direct protein sequencing; Immu DOMAIN 24 34 CO DOMAIN 35 49 FT DOMAIN 50 56 CO DOMAIN 50 56 CO DOMAIN 89 97 CO DOMAIN 89 108 FT DOMAIN 98 108 BY NON TER 108 108 BY NON TER 108 108 BY
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21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
1g kappa chain V-V region HP 123E6.
Mus musculus (Mouse)
Eukarvor: ...
      Direct protein
DOMAIN
JOMAIN
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                                                                                              Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                             "Complete amino acid sequence of light chain variable reg
from five monoclonal anti-p-azophenylarsonate antibodies
with respect to a crossreactive idiotype.";
Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
-!- MISCELLANEOUS: Anti-arsonate hybridoma protein.
HSSP, P01594; 1JV5.
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STRAIN=A/J;
MEDLINE=82150934; PubMed=6801658;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                      InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Siegelman M., Capra J.D.;
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Rodentia;
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Framework-1.
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Pred. No. 1.
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By similarit
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Complementarity-determining-3.
Complementarity-determining-1
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21-JUL-1986 (Rel.
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Ig kappa chain V-V
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P01648;
                                                                                                          DISULFID NON_TER
                                                                                                                                                                                                                                                        with respect to a crossreactive idiotype."; Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
                                                                                                                                                                                                                                                                               "Complete amino
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                        STRAIN=A/J;
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                                                                         Score 477;
Pred. No. 9.
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Framework-3.
Complementarity-determining-3.
Framework-4.
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X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Klausner R.D., Collins F.S., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang L., Scheetz T.E.,

X Hopkins R.F., Jordan H., Moore T., Abramson R.D., Mullahy S.J.,

X Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

X Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

X Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

X Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunaratne P.H.,

X Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

X Hilalon D.K., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

X Hilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

X Halkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

X Halkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

X Hones S. I. Malska U., Smailus D.E., Schnerch A., Schein J.E.,

X Jones S. I. Malska U., Smailus D.E., Schnerch A., Schein J.E.,
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01-DEC-2001
01-MAR-2004
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Pfam; PF00047; 1g; 1.
SMART; SM00406; IGV; 1.
SROSITE; PS50935; IG LIKE; 2.
PROSITE; PS00290; IG MHC; UNKNOWN 1.
PROSITE; PS00962; RIBOSOMAL_S2_1; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jones S.J., Marra M.A.;
Jones S.J., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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SEQUENCE 2
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STRAIN=FVB/N; TISSUE=Colon;
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STRAIN=FVB/N; TISSUE=Colon;
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terPro; IPR007110; Ig-like.
terPro; IPR003597; Ig-q1.
terPro; IPR003006; Ig_MCC.
terPro; IPR003596; Ig_v.
terPro; IPR003596; Ig_v.
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                                             RFSGGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
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Pred. No. 2.9e
6; Mismatches
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2.9e-39;
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RESULT 12
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ID KV5U MOUSE
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DT 13-AUG-1987 (
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Q9JL84;
01-OCT-2
                            This SWISS-PROT entry is copyright. It is produbetween the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There a use by non-profit institutions as long as i
                                                                                                                                                                       MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Mil
"mRNA sequences define an unusually restricted
phenyloxazolone and its early diversification."
                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
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EMBL; AF206022; AAF69320.1; -.
HSSP; P01594; LJV5.
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Malkiel S., Liao L., Cunningham M.W., I
"T-Cell-dependent antibody response to
          modified
                                                                                                                                Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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Mammalia; Eutheria;
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(TremBLrel. 15, Last sequence up
(TremBLrel. 25, Last annotation
immunoglobulin light chain varia
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Sciurognathi; Muridae; Murinae; Mus
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02-MAR-2004 (TrEMBLrel. 2:
02-MAR-2004 (TrEMBLrel. 2:
02-MAR-2004 (TrEMBLrel. 2:
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InterPro; IPR003596; Ig_v.
Pfam; PP00047; Ig; 1.
SWART; SW00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Hybridoma; Immunoglobulin V re;
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DOMAIN
                                                                                                                                                                   STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.,
"Antinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY436892; AAR11052.1; -.
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             GGGSGTDYSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLE
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Sciurognathi; Muridae; Murinae;
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02-MAR-2004 (TrEMBLrel. 27, Created)
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Liang Z., Xie C., Chen C., Kreska D., Hsu K.,
"Antinuclear autoantibodies from B6.Sle1 mice.
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ
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ANA immunoglobulin kappa light chain
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Mammalia; Eutheria;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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                     YSLTISNLEPEDIATYFCHQYSKLPWTFGGGTKLEIKR
                                                                                                            LPASLGDRVTISCSASQDISNYLNWYQQKPDGTVKLLIFYSSNLHSGVPSRFSGGGSGTD
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FSLTISNLDQEDIATYFCQQGNS:
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11250 MW;
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82.8%;
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83.7%;
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from B6.Sle1 mice.
EMBL/GenBank/DDBJ
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ght chain (Fragment).
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	0	տ	4	w	2	۳	Result No.
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71.4	71.4	٠	71.5		71.7	71.7	71.7	71.7	71.7	71.8	71.8	71.9	72.2	72.2	72.2	83.4	84.4	88.4	•	٠	٩	89.9	٠	100.0	Query Match
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Query Match 100.0%; Score 3071; DB 4; Length 582; Best Local Similarity 100.0%; Pred. No. 4.3e-149; Matches 582; Conservative 0; Mismatches 0; Indels 0

0

Gaps

0

SQ Sequence 582 AA;	The present invention describes a monoclonal antibotic specifically with ganglioside GD3. The antibody and CC useful in the diagnosis and therapy of tumours, par CC diagnosis. The present sequence is a protein used it compared to the invention	PS Claim 39; Page 175-179; 183pp; Japanese.	New human type complementation-determining region- and derivatives against ganglioside GD3, useful in of e.g. tumors, with low antigenicity, little side activity in cancer.	DR WPI; 2001-266143/27.	PI Hanai N, Shitara K, Nakamura K, Niwa R;	PA (KYOW) KYOWA HAKKO KOGYO KK. XX	PR 30-SEP-1999; 99JP-00278291. PR 06-APR-2000; 2000JP-00105088.	PF 29-SEP-2000; 2000WO-JP006774.	PD 05-APR-2001. XX	PN WO200123432-A1.	OS Synthetic. XX	KW Ganglioside; GD3; complementarity determining region; KW cancer.	Ganglioside GD3 specific antibody related protein	DT 03-JUL-2001 (first entry)	AC AAB81991;	RESULT 1 AAB81991 ID AAB81991 standard; protein; 582 AA.	ALIGNMENTS	26 2191.5 71.4 447 6 AAE33524 27 2189.5 71.3 463 8 ADM72025 28 2184.5 71.2 451 8 ADB84594 29 2184.5 71.1 446 7 ADF11425 30 2184 71.1 449 3 AAY68810 31 2181 71.0 449 3 AAY68810 32 2181 71.0 474 6 ABU00017 34 2181 71.0 474 7 ADF65775 36 2180.5 71.0 461 6 ABR39947 37 2180.5 71.0 461 6 ABR39943 38 2177.5 70.9 442 6 ABB0109 40 2177.5 70.9 442 6 ABB0109 41 2177.5 70.9 442 6 ABB0109 42 2177.5 70.9 442 6 ABB0109 43 2177.5 70.9 442 6 ABB0109 44 2177.5 70.9 442 6 ABB0109 45 2177.5 70.9 442 7 ADE94066 46 2177.5 70.9 461 6 ABR39848 47 2177.5 70.9 461 6 ABR39848
	noclonal antibody which can react to antibody and its derivatives are if tumours, particularly cancer protein used in the exemplification		transplanted antibody diagnosis and therapy effects but potent									n; CDR; antibody;	SEQ ID NO: 57.					Aae33524 Human AQC Adm72025 Chimeric Adm13584 008 heavy Adf11425 2D8 anti- Adp88494 Humanised Aay68810 A rat hea Aau014065 Heavy cha Abu08017 Human mon Adf65775 Human mon Adf65775 Human soJ Abr39847 Hu266 N56 Abr39843 Hu266 N56 Abr39843 Hu266 Humanised Abu60109 Heavy cha Adde94066 Humanised Abu60719 Heavised Adm61714 Humanised Adm61746 Humanised Adm61748 Hu266 N56

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                                                                                                                                                                                     Ganglioside; GD3; complementarity
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                                     Hanai N,
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          Unidentified
                           Ganglioside;
                                                               02-JUL-2001
                                                                                   AAB83156;
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                             GM2;
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FFFXSXSSSSSXS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 582 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 41; Page 168-172; 183pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                     LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSSA
                                                                                                                                                                                                           TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                                                                                                               TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                                                                                                                                                                TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
                                                                                                                                                                                                                                                                                                    TYRVVSVLTVLHQDWLNGKEYKCKVSNKALÞAÞÍEKTISKAKGQPREÞQVYTLÞÞSRDEL
                                                                                                                                                                                                                                                                                                                                                      SVFLPPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                                                                                                                                                                                                        SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                                                                                                                                                                                                                                                          LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSSA
IVLELKGSETTFMCBYADETATIVEFLNRWITFCQSIISTLT
                       IVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 582
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98.3%;
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Pred. No. 8.7e-147;
5; Mismatches 5;
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AB83156

AAB83156 standard; protein; 583 AA.

C AAB83156;

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R 02-JUL-2001 (first entry)

K S Ganglioside GM2 antibody-related protein
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antibody; cytostatic; cytotoxic;

cancer.

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RESULT 4
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Best Local Simi
Matches 523;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to derivatives of an antibody against ganglioside GM2. The antibody may be a monoclonal antibody or its fragments. The antibody is combined with a radioactive isotope, protein or small drug in the treatment and diagnosis of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monoclonal antibodies against ganglioside GM2 combined with drugs, radioisotopes or proteins for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hanai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KYOW ) KYOWA HAKKO KOGYO KK.
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                                                                                                                                                                                                                                                                                                                                                                                           1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                               SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQ
                                                                                                                                                                                                                                                                                                                                                     SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYY---FDSWGQGTTLTV
                                                                                 NNYKOPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISN
                                                                                                                      RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGI
                                                                                                                                                           DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                                              YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                                                                                   GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                                                                                                                                                                                                           SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
                                                                                                                                                                                                                                                                       SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
                                                                                                                                                                                                                                                                                                                                     NQKFKSKVTITVDTSTSTAYMELHSLRSEDTAVYYC--ATYGHYYGYMFAYWGQGTLVTV
                                                                                                                                                                                                                                                                                                                                                                         583 AA;
                                 INVIVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT
                                          INVIVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 582
                                                                                                           RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGI
                                                                                                                                                DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                                   YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                                                                         GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                       NNYKNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNPHLRPRDLISN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 61-65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nakamura K,
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                       89.9%;
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Pred. No. 3.6
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3.6e-133;
nes 36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New recombinant anti-EpCAM antibody having an amino acid sequence defining an immunoglobulin light or heavy chain framework region, for the diagnosis, prognosis and treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
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STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                 PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                                                                                                                                                    GLYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                                                                                                                                                                                     ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                                                                                                                                                                                          ADDFKGRFAFSLETSTSTAFLQINNLRSEDTATVFCVRFISKGDV----WGQGTSVTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89.4%; Score 2744.5; DB 6; ilarity 88.5%; Pred. No. 2.1e-132; Conservative 31; Mismatches 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 80-82; 82pp; English.
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                                                                       Query Match
Best Local S
Matches 515
                                                                                                                                                                                                The invention relates to cytokine fusion proteins with increased therapeutic index and methods for increasing the therapeutic index of such fusion proteins. The fusion protein comprises a non-interleukin-2 (IL-2) moiety fused to a mutant IL-2 moiety. It is useful for preparing a composition for treating cancer, viral infections or immune disorders. The fusion protein is also used in gene therapy. The present sequence is dI-KS-ala-IL2 (D20T) variant protein comprising dI-KS heavy chain fused to human IL-2 (D20T) variant protein. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New fusion protein comprising a non-IL-2 molety fused to a mutant IL-2 molety, useful for preparing a composition for treating cancer, viral infections or immune disorders.
                                                                                                                                                         Sequence 579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gillies
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12-APR-2002; 2002US-0371966P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2002; 2002WO-US038780
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10;
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                                                                  Local Simphes 515;
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Human type complementation-determining domain transplanted antibody and derivatives against ganglioside GD2, useful in diagnosis and therapy of e.g. tumors, has low antigenicity, little side effects but potent activity in cancer.

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RESULT 7
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Matches 518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes an antibody, which can react specifically with ganglioside GD2, and is transplanted with a human type complementation-determining domain (CDD) or its fragments. The antibody and its derivatives are useful in diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
                       WO2004055056-A1
                                                                                                      immunoglobulin; variace cancer; cell surface
                                                                                                                                                                Humanised immunoglobulin heavy chain-IL-2
                                                                                                                                                                                                                                                                                            ADP42961
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGINNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGINNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISNINV
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                                                                                                                         variable region; antibody;
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                                                                                                                                                                                                                                                                                            protein;
                                                                                                   glycosphingolipid;
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Pred. No. 9.7e-132;
4; Mismatches 39;
                                                                                                                                                                  fusion
                                                                                                                                                              protein SEQ ID NO:6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibody variable region specifically binds to human cell surface glycosphingolipid GD2. An antibody variable region of the invention cytostatic activity, and may have a use in gene therapy. The antibod be used for treating cancer. The nucleic acid or cell is useful for manufacturing a medicament that may be used for treating diseases su cancer. The present sequence represents an immunoglobulin heavy chail 2 fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New modified m14.18 antibodies with reduced immunogenicity specifically bind the human cell surface glycosphingolipid for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local 517;
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DB; ADP42959.
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                                                                                    QGNVFSCSVMHEALHNHYTQKSLSLSPGKAPTSSSTKKTQLQLEHLLLDLQMILNGINNY
                                                                                                                                                                             TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
                                                                                                                                                                                                                                                                       TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL
                                                                                                                                                                                                                                                                                                                                                              SVFLFPPKPKDTLMI SRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                                                                                                                                                                                                                                                                                                                                                                  LYSLSSVYTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NQKFKGRATLTVDKSTSTAYMHLKSLRSEDTAVYYCVS---GMEY---WGQGTSVTVSSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTFAKRLEWVAYISSGGSGTYY
KNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISNINV
                                                          QGNVFSCSVMHEALHNHYTQKSLSLSPG-APTSSSTKKTQLQLEHLLLDLQMILNGINNY
                                                                                                                                                    TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
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88.8%; Pred. No. 6.3e-131;
tive 24; Mismatches 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                              Matches 494;
                                                                                                                                                                                                                                      The invention relates to cytokine fusion proteins with increased therapeutic index and methods for increasing the therapeutic index of such fusion proteins. The fusion protein comprises a non-interleukin-2 (IL-2) moiety fused to a mutant IL-2 moiety. It is useful for preparing a composition for treating cancer, viral infections or immune disorders. The fusion protein is also used in gene therapy. The present sequence is Gl-NHS76 (gamma4h) (FN>AQ)-ala-IL2 (D2OT) variant protein comprising dI-KS heavy chain fused to human IL-2 (D2OT) variant protein. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                     Sequence 580 AA;
                                                                                                                                                                                                                                                                                                                                        Example 10;
                                                                                                                                                                                                                                                                                                                                                           New fusion protein comprising a non-IL-2 momoiety, useful for preparing a composition infections or immune disorders.
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12-APR-2002; 2002US-0371966P.
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                 GLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVDKRVEPKSCDKTHTCPPCPAPEFLGG
                             GLYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                           ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                                          EVILVESGGDFVKPGGSLKVSCAASGFAFSH-YAMSWVRQTPAKRLEWVAYISSGGSGTY
                                                          ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                YNPSLKSRVTISVDTSKNQFSLKLSSVTAADTAVYYCARGKWSK--FDYWGQGTLVTVSS
                                                                                                                   YSDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
                                                                                                                                       QVQLQESGPGLVKPSETLSLTCAVSGYSISSGYYWGWIRQPPGKGLEWIGSIYHSGS-TY
PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (gamma4h)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interleukin-2; IL-2; cancer; viral infection; immune disorder;
apy; immunoglobulin; Ig; fusion protein; human; variant.
                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                         Page 68-71;
                                                                                                                                                                                                                                                                                                                                                                                                                                          LEXIGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (FN>AQ)-ala-IL2
                                                                                                                                                                                       84.4%;
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                                                                                                                                                                                                                                                                                                                                       71pp;
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                                                                                                                                                                              Score 2592; DB 6;
Pred. No. 1.3e-124;
1; Mismatches 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (D20T) variant
                                                                                                                                                                                                                                                                                                                                                                    moiety fused to on for treating c
                                                                                                                                                                              54;
                                                                                                                                                                                              Length 580;
                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                    o a mutant IL-2 cancer, viral
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RESULT 9
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                                     The invention relates to cytokine fusion proteins with increased therapeutic index and methods for increasing the therapeutic index of such fusion proteins. The fusion protein comprises a non-interleukin-2 (IL-2) molety fused to a mutant IL-2 molety. It is useful for preparing a composition for treating cancer, viral infections or immune disorders. The fusion protein is also used in gene therapy. The present sequence is dI-NHS76 (gamma2h) (FN>AQ)-ala-IL2 (D2OT) variant protein comprising dI-KS heavy chain fused to human IL-2 (D2OT) variant protein. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-DEC-2001;
12-APR-2002;
                                                                                                                                                                                                                                                          Claim 37;
                                                                                                                                                                                                                                                                                                New fusion protein comprising a non-IL-2 moiety fused to a mutant IL-moiety, useful for preparing a composition for treating cancer, viral infections or immune disorders.
                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-513757/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-DEC-2002;
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Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQAQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VIVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QQGNIFSCSVMHEALHNHYTQKSATATPGAAPTSSSTKKTQLQLEHLLLTLQMILNGINN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STYRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YKNPKLTRMLTFKFYMPKKATELKHLQCLEEELKPLEEVLNLAQSKNFHLRPRDLISNIN
                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interleukin-2; IL-2; cancer; viral infection; immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LEXIGEN RES
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2002US-0371966P.
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                                                                                                                                                                                                                                                        71pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CENT CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          582
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RESULT 10
AAO18400
ID AAO18
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AC AAO18
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XX Mouse
KW Mouse
KW neopl
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XX WO200
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XX I3-00
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Best Local Simi:
Matches 491; (
                      13-OCT-2000; 2000US-0240285P
13-MAR-2001; 2001US-0275289P
21-JUN-2001; 2001US-0299987P
                                                                                                                                                                     Mouse; humanised antibody; lymphotoxin beta receptor; antibody; cancer; neoplasia; LT-beta-R; light chain; heavy chain; variable region.
                                                                                                                                                                                                            Mature
                                                                                                                                                                                                                                  11-OCT-2002
                                                                                                                                                                                                                                                                               AAO18400 standard;
                                                                    12-OCT-2001; 2001WO-US032140
                                                                                            18-APR-2002
                                                                                                                  WO200230986-A2
   (BIOJ ) BIOGEN INC
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; Pred. No. 5.6e-123;
31; Mismatches 56;
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Best Local Similarity
Matches 418; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to humanised anti-lymphotoxin beta receptor (LT-beta-R) antibodies. These are derived from the murine LT-beta-R binding antibody CBE11 and can be used to treat neoplasia in humans. The present sequence is a humanised murine CBE11 heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New humanized anti-lymphotoxin-beta receptor antibody, useful for treating or reducing the advancement, severity or effects of neoplasia, particularly solid tumors (i.e. carcinomas) including colorectal cancer and breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5; Page 25-26; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Garber
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'n
             QQGNVFSCSVMHEALHNHYTQKSLSLSPG 448
                                                   LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
                                                                    LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
                                                                                                                            STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                                                PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
                                                                                                                                                                                                                      GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                                                                                                                                                                       GLYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
                                                                                                                                                                                                                                                                               ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                                                                                                                                                                                                                     PDSVKGRFTISRDNAKNSLYLQMSSLRAEDTAVYYCAREENGNFYYFDYWGQGTTVTVSS
                                                                                                                                                                                                                                                                                                                                                   SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGT-YYFDSWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                           EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
QQGNVFSCSVMHEALHNHYTQKSLSLSPG
                                                                                                          STYRVVSVLTVLHQDWLNGKBYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
                                                                                                                                                                                                                                                                                                                                                                                            EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMYWFRQAPGKGLEWVATISDGGSYTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                 72.2%;
nilarity 93.1%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 2218.5; DB 5;
Pred. No. 1.2e-105;
9; Mismatches 21;
449
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ADN97491
ID ADN97491;
XX ADN974
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RESULT 12
ADN97489
ID ADN97
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AC ADN97
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DT 01-JU
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Best Local Sim
Matches 419;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an artificial proprotein comprising three peptide sequences: a first peptide sequence of interest, a propeptide sequence attrached to the C-terminus of the first peptide sequence of interest, and a second peptide of interest attached to the C-terminus of the propeptide sequence. The artificial proprotein and polynucleotides are useful for artificial multimeric protein engineering, e.g. antibodies and antibody fragments in eukaryotes. This sequence corresponds to a protein used in the generation of the protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New artificial proprotein comprises three peptide sequences, useful artificial multimeric protein engineering in eukaryotes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-OCT-2003; 2003WO-US031420
            01-JUL-2004
                                 ADN97489;
                                                      ADN97489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LARG-) LARGE SCALE BIOLOGY CORP
                                                                                                                            413
                                                                                                                                                 617
                                                                                                                                                                      353
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                                                                                                                                                                                                                                                                             437
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                                                                                                                                                                                                                                                                                                                                            113
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DB; ADN97490.
                                                                                                          677
                                                                                                                                                                                                               293
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                                                                                                                                                                                                                                                                                                                                                                                                                             <u>بــر</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; SEQ ID NO 64; 244pp; English.
                                                                                                                    TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                     PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYT
                                                                                                                                                                                                                                                 APELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTK
                                                                                                                                                                                                                                                                                                                                                                                    SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKL-----GTYYF---DSWGQG
                                                     standard;
                                                                                                                                                              LPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKL
                                                                                                                                                                                                                                                                                           PAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCP
                                                                                                                                                                                                                                                                                                                       ASVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTF
                                                                                                                                                                                                                                                                                                                                      TTLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTF
                                                                                                                                                                                                                                                                                                                                                                PDSVKGRFTISRDNDKNALYLQMNSLKSEDTAMYYCARRSEFYYYGNTYYYSAMDYWGQG
                                                                                                                                                                                                                                                                                                                                                                                                         EVDLVESGGDLVKPGGSLKLSCAASGFTFSHYGMSWVRQTPDKRLEWVATIGSRGTYTHY
                                                                                                                                                                                                                                                                                                                                                                                                                     EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            713 AA;
                                                                                                      TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                 LPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKL
                                                                                                                                                                                          PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYT
                                                                                                                                                                                                                                      APELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKPNWYVDGVEVHNAKTK
                                                                                                                                                                                                                                                                             PAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Edwards P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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            (first
                                                     protein;
            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            72.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  10;
                                                     715
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 2216; DI
Pred. No. 2.6e
.0; Mismatches
                                                     A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 713;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                       419;
                                                                                                                                                                                                                                                                                                                                                                                               715 AA;
                                                                                                                                                                                                                                                                                                                                                       Conservative
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New artificial proprotein comprises three peptide sequences, artificial multimeric protein engineering in eukaryotes.
                                                                                                                                                                                                                                                     03-OCT-2002; 2002US-0415940P
                                                                                                                                                                                                                                                                                                    03-OCT-2003; 2003WO-US031420.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      artificial proprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Artificial protein construction protein #3
                                                                                                                                                                                                                                                                                                                                                                                                     WO2004031362-A2
                                                                                                                                                                                                      (LARG-) LARGE SCALE BIOLOGY CORP
                                                                         2004-330170/30.
DB; ADN97488.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        propeptide; protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      engineering;
                            useful
                               for
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peptide sequences: a first peptide sequence of interest, a propeptide sequence attached to the C-terminus of the first peptide sequence of interest, and a second peptide of interest attached to the C-terminus of the propeptide sequence. The artificial proprotein and polymucleotides are useful for artificial multimeric protein engineering, e.g. antibodies and antibody fragments in eukaryotes. This sequence corresponds to a protein used in the generation of the protein of the invention. The invention relates to an artificial proprotein comprising three peptide sequences: a first peptide sequence of interest, a propeptide SEQ ID ö 62; 244pp; English.

72.2%;

Score 2216; Pred. No. 2. Mismatches

1.6e-105; 1es 20; BB

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8

Length Indels

10;

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PAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYPPEPVTVSWNSGALTSGVHTF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SDSVKGRFTISRDNAKWTLYLQMRSLRSEDSAMYFCTRVKL-----GTYYF---DSWGQG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                PDSVKGRFTISRDNDKNALYLOMNSLKSEDTAMYYCARRSEFYYYGNTYYYSAMDYWGQG
TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                              TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                           PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYT
                                                                                                                                                                                                             PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKV$NKALPAPIEKTISKAKGQPREPQVYT
                                                                                                                                                                                                                                                                                         APELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTK
                                                                                                                                                                                                                                                                                                                                                        PAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                ASVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVDLVESGGDLVKPGGSLKLSCAASGFTFSHYGMSWVRQTPDKRLEWVATIGSRGTYTHY
  715
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                                                                                          678
                                                                                                                                      412
                                                                                                                                                                               618
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                                                                                                                                                                                                                                                                                                                                                            498
                                                                                                                                                                                                                                                                                                                                                                                                      232
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RESULT 13
ADM47075
ID ADM477075
AC ADM477
XX ADM477
XX ADM477
XX MOUSE
XX MOUSE
XX MOUSE
XX MOUSE
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                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the production of a methylotroph yeast that produces mammalian sugar chains, comprising disrupting the OCH1 gene in the yeast that encodes for alpha-1,6-mannosyl transferase and inserting and expressing the alpha-1,2-mannosyl transferase and inserting cand expressing the alpha-1,2-mannosidase gene. The specification also includes DNA sequences encoding: (a) orcidin-5'-phosphate decarboxylase (URA3); (b) phosphoribosyl-amino-imidazole succinocarboxamide synthase (ADE1); (c) imidazole-glycerol-phosphate dehydratese (HIS3); (d) 3-cisopropyl malate dehydrogenase (LEU2); (e) alpha-1,6-mannosyl transferase (OCH1); (f) proteinase A (PE4); (g) proteinase B (PRE1); and (h) aspartic protease (YPS1), mannosyl transferase (KTR1 or MNN9), alcohol oxidase (AOX) and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) gene sequences. The yeast is used for the production of human and mammalian high mannose glycoproteins with high yield and purity. The method is also useful for producing hybrid or complex sugar chains containing mammalian type chains. This sequence represents a mouse anti-human G-CSF antibody theavy chain used in the invention
                                                                                                                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Producing methylotroph yeast disrupting the OCH1 gene and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6-mannosyl transferase; alpha-1; 2-mannosidase; orotidin-5'-phosphate decarboxylase; URA3; phosphoribosyl-amino-imidazole succinocarboxamide synthase; ADE1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 28; SEQ ID NO 94; 247pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-APR-2003; 2003WO-JP005464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       imidazole-glycerol-phosphate dehydratase; HIS3;
3-isopropyl malate dehydrogenase; LEU2; proteinase A; proteinase B;
PEP4; YPS1; KTR1; MNN9; AOX; GAPDH; mannosyl transferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            methylotroph yeast; mammalian sugar chain; OCH16-mannosyl transferase; alpha-1; 2-mannosidase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM47075 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KIRI ) KIRIN BEER KK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUN-2004 (first entry)
   80
                                                                                                                                                                                                                                          416;
                                                                                                                                                                                                                                                                                                                                                                                                                               chain used
                                                         61
                                                                                                                        20
                                                                                                                                                                              μ
                                                                                                                                                                                                                                                                             Similarity
                               EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                   475 AA;
                                                                                                                        EVQLVESGGGLVKPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWVSSISSSSYIYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>~</u>
                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kitagawa Y,
                                                                                                                                                                                                                                                                       71.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Komeda T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             that expresses mammalian sugar chains by inserting an alpha-1,2-mannosidase gene.
                                                                                                                                                                                                                                                                       Score 2209.5;
Pred. No. 3.76
                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₿
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                                                                                                                                                                                                                                                                          .7e-105;
                                                                                                                                                                                                                                                                                                      DB 7;
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                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                   Length 475;
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ARAFOTAT 14
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ID ARAFOTAS3
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AC AARAFOTAS3
AC CHime
KW CHITE
AC COLOTY
AC COLOTY
AC COLOTY
AC The B
CC murin
CC carci
             The sequences (AAQ54651-52) show the light and heavy chain cDNAs of murine T84.12. The T84.12 antibody is directed against the tumour mac carcinoma embryonic antigen, and is useful for tumour imaging and immunotherapy. The amino acid sequence given in the specification habeen incorrectly identified as a nucleic acid sequence, therefore unacceptable characters have been represented as an 'N'. The amino a sequence given below has been derived from the cDNA, by the indexer. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                              New chimeric T 84.12 antibody active against carcinoembryonic antigen - has murine variable and human constant regions, also DNA encoding it and transformed myeloma cells.
                                                                                                                                                                                                                                                                                                     WPI; 1994-007204/01.
                                                                                                                                                                                                                                                                                                                                      Shively JE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric; carcinoembryonic antigen; CCA; murine; mouse; constant; region; transform; myeloma cell; light chain; tumour.
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24-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                            (YANG/)
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                                                                                                                                                                            1; Page 22-23;
                                                                                                                                                                                                                                                                                                                                                                          YANG
CITY
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Best Local
                                                                                                                                                                                                                                                                                                                                                                 Mus sp.
Antibody against the N terminus of glypican 3(GPC3) disruption and is useful as an anticancer agent.
                                                            WPI; 2004-269573/25.
N-PSDB; ADM72028.
                                                                                                                                                                                                                                                                                                                                                                                                                                               GPC3; glypican 3; anti-GPC3 antibody; cell disruption; anti-cancer; cytostatic; M19B11.
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                                                                                                                                                                                                    04-SEP-2002; 2002WO-JP008999
                                                                                                                                                                                                                                            04-SEP-2003; 2003WO-JP011318
                                                                                                                                                                                                                                                                                    18-MAR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric mouse-human antibody M19B11 heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JUN-2004
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                                                                                                                         Aburatani
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Pred. No. 6.3e-105;
8; Mismatches 18;
                                                                                                                     Nakano
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                                                                                                                     Ohizumi
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                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 71.8%;
Best Local Similarity 92.3%;
Matches 418; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an antibody against the N terminus of glypican 3 (GPC3). The antiboduy can be used for causing cell disruption and can be uses as an anti-cancer agent. The present sequence represents a chimeric mouse-human antibody M19B11 heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 471 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; SEQ ID NO 14; 122pp; Japanese
                                                   379
439
                       417
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                                                                                                                                                      LGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREE
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             SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                                                                         QSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCBAPEL
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                                                                                                                                                                                                                                                                                                                                                                 EVQLVESGGDLVKPGGTLKLSCAASGSTFSNYAMSWVRQTPEKRLEWVAAIDSNGGTTYY
                                                  RDELTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVDK
                                                                 RDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDK
                                                                                                     QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPS
                                                                                                                    QYNSTYRVVSVLTVLHQDWIMGKEYKCKVSNKALFAPIEKTISKAKGQFREPQVYTLFPS
 SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                           QSSGLYSLSSVVTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPEL
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 2203.5;
Pred. No. 7.5e.
9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                      9;
                                                                                                                                                                                                                                                                                                                                                                                                                      5; DB 8;
7.5e-105;
hes 21;
                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Search completed: December 23, 2004, 18:57:43
Job time: 157.692 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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are greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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Match
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3071
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    IBSUEd_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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US-09-234-340A-71
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US-08-466-153B-8
US-08-867-352B-14
US-08-887-352B-16
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US-09-109-207C-16
US-09-109-207C-16
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US-09-110-207C-16
US-09-111-14
US-09-216-025-14
US-09-216-025-16
US-09-111-966-14
US-09-113-996-14
US-09-113-996-14
US-09-109-207C-18
US-09-109-207C-18
US-09-109-207C-18
US-09-109-207C-18
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US-09-26-005-18
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Sequence 7, Appli
Sequence 71, Appl
Sequence 71, Appl
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RESULT 2
US-09-027-449-71
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Patent No. 6
                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/027,449

FILING DATE: 20-Feb-1998

CLASSIFICATION: 435
                                                  APPLICATION NUMBER: 60/074,330
PRIOR APPLICATION DATA: 2-Jan-1998
PRIOR APPLICATION DATA: 60/038,664
APPLICATION NUMBER: 60/038,664
FILING DATE: 21-Feb-1997
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 34,659
REGISTRATION NUMBER: 34,659
                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Gonzal
REFERENCE/DOCKET NUMBER: 91(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates and
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibodies
NUMBER OF SEQUENCES: 72
                                                                                                                                                                                          PRIOR APPLICATION DATA:
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Matches 401;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 452 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                       RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                         DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                                                                                                                                                                                                         SSGLYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
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RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                   EEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                                                YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                                        YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                                                                                                    GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                                                                                                                                                                                                                                                                                 SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EVQLVQSGGGLVQPGGSLRLSCAASGYSFSSHYMHWVRQAPGKGLEWVGYIDPSNGETTY
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88.7%;
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Pred. No. 2.66
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GENERAL INFORMATION:
APPLICANT: Gonzalez, Tania R.
APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fr:
TITLE OF INVENTION: Antibody Fr:
TITLE OF INVENTION: Humanized Ar
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 71, Appl
Patent No. 613342
                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb fl.
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winpatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/99/026,985
FILING DATE: 20-Feb-1998
                                                                                                                                                                                                                             STREET: 1 CITY: South San Fra
CITY: South San Fra
CTATE: California
CLASSIFICATION:
ATTORNEY/AGENT IN
                                                                                                                                                                                                                ZIP: 94080
                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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1 DNA Way
San Francisco
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      INFORMATION:
                                                                US/09/026,985
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d Anti-IL-8 Monoclonal Antibodies
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RESULT 4
US-09-121-952A-71
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TELEPHONE: 650/225-5530
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
LENGTH: 452 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                              Sequence 71, Application US/09121952A Patent No. 6458355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 70.0%;
Best Local Similarity 88.7%;
Matches 401; Conservative 1
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APPLICANT: Genent
APPLICANT: Koumen:
APPLICANT: Leong,
                                                                                                                                       APPLICANT: Leong, Steven R.

APPLICANT: Presta, Leonard G.
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
TITLE OF INVENTION: METHODS OF TREATING INFLAMMATORY DISEASES
TITLE OF INVENTION: MITH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44
COMPUTER: IBM PC compatible
                                                                STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
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Koumenis, Iphigenia
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                                                                                                                                Inc.
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 452 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WINDATA: (Genentech) CURRENT APPLICATION DATA: APPLICATION UMBER: US/09/121,9: FILING DATE: 24-Jul-1998 CLASSIFICATION DATA: APPLICATION UMBER: 60/074330 PILING DATE: 22-JAN-1998 PRIOR APPLICATION DATA: APPLICATION UMBER: 60/074330 PILING DATE: 22-JAN-1998 PRIOR APPLICATION UMBER: 60/075467 PILING DATE: 20-FEB-1998 ATTORNEY/AGENT INFORMATION: NAME: LOVE, Richard B. 1660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P108
TELECOMPUNICATION INFORMATION:
TELEPHONE: 650/225-5530
TELEPAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 452 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                          361
421
                  418 RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                   358 DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
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                                                                          EEMTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVDKS
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88.7%;
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Pred. No. 2.6e-159;
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452
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RESULT 5
US-09-234-340A-71
Sequence 71, Application US/09234340A
Patent No. 6468532
GENERAL INFORMATION:
APPLICANT: Genentech, Inc., Heei, Va.
APPLICANT: Koumenis, Iphigenia
APPLICANT: Leong, Steven R.
APPLICANT: Preeta, Leonard G.
APPLICANT: Shahrokh, Zahra

360 417 237 240 297 297 300 357

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US-09-234-340A-71
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Best Local (
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US/09/
PILING DATE: 24-Jul-1998
APPLICATION NUMBER: 60/074
PILING DATE: 22-JAN-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/0:
FILING DATE: 20-FEB-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/234,340A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zapata, G
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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California
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EEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                   DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
                                                                                                                                      GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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                                                                   YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                          YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
                                                                                                                                                                      GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ 297
                                                                                                                                                                                                          SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
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1 DNA Way
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METHODS OF TREATING INFLAMMATORY DISEASES

WITH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES:

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Pred. No. 2.6e-159;
9; Mismatches 29;
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US-08-466-151-8
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APPLICANT: Jardieu, E
APPLICANT: Presta, Le
TITLE OF INVENTION: 1
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/08466151 Patent No. 6037453
                                                                                                                                                                                                                                                                                                                                                                          Matches 403;
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: 3.5 inch, 1.44 Mb flucomputer: IBM PC compatible operating system: PC CDS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION NUMBER: US/08/466,151 FILING DATE: CLASSIFICATION DATA: PRICING DATE: OB/466163 APPLICATION NUMBER: 08/466163 APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: 08/466163 APPLICATION DATA: APPLICATION DAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 453 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/744
PILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/405
FILING DATS: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185
FILING DATS: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
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CITY: S
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SSAST--KGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
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Pred. No. 2.6e-157;
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PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR FILING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR FILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 64
SEQ ID NO 8
SEQ ID NO 8
SEQ ID NOS: 64
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US-08-466-163B-8
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Best Local S
Matches 403
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APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
FILE REFERENCE: P0718P2C1D1
CURRENT APPLICATION NUMBER: US/08/466,163B
CURRENT FILING DATE: 1995-06-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial sequence FEATURE:
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LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWXVDGVEVHNAKTKPRE
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                                                                                 LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
                                                                                                                                                                       SSAST - - KGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
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Pred. No. 2.6e
21; Mismatches
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APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Preventing th
FILE REFERENCE: P0718P2C3US
CURRENT APPLICATION NUMBER: US/09/802,096
CURRENT FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR FILING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR FILING DATE: 1992-08-14
PRIOR FILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR PILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
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US-09-802-096-8
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SEQ ID NO 8
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Best Local Similarity
Matches 403; Conserva
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TYPE: PRT
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                                            EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
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                     EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
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KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

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APPLICANT: JARTHER, PAULA M.

APPLICANT: Presta, Leonard G.

TITLE OF INVENTION: Method of Treating Allergic Dis

FILE REFERENCE: P0718P2C2US

CURRENT APPLICATION NUMBER: US/09/802,077

CURRENT APPLICATION NUMBER: US 08/405,617

PRIOR APPLICATION NUMBER: US 08/405,617

PRIOR FILING DATE: 1995-03-15

PRIOR FILING DATE: 1995-03-15

PRIOR APPLICATION NUMBER: US 08/185,899

PRIOR FILING DATE: 1992-01-26

PRIOR FILING DATE: 1992-08-14

PRIOR FILING DATE: 1992-08-14

PRIOR APPLICATION NUMBER: US 07/879,495

PRIOR APPLICATION NUMBER: US 07/744,768

PRIOR APPLICATION NUMBER: US 07/744,768

PRIOR FILING DATE: 1991-08-14

NUMBER OF SEQ ID NOS: 64

I FRIOR FILING DATE: 1991-08-14
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US-09-802-077-8
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Best Local &
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Similarity 88.8%; Pr
03; Conservative 21;
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KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                  SREEMTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTVD
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Pred. No. 2.6e-157;
Pred. No. 2.6e-24;
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RESULT 10

US-08-887-352B-14
; Sequence 14, Application US/08887352B
; Patent No. 5994511
; GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta ITILE OF INVENTION: Improved Anti-IgE Antibod ITILE OF INVENTION: Improving Polypeptides NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb flo

MEDIUM TYPE: 3.5 inch, 1.44 Mb flo

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOTTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/887,352E

FILING DATE: 03-Jul-1997

CLASSIFICATION: 530

ATTORNEY/ACENT INFORMATION:

NAME: Svoboda, Craig G.

REGERENCE/DOKET NUMBER: 39,044

REFERENCE/DOKET NUMBER: 39,044

REFERENCE/DOKET NUMBER: 37,044

REFERENCE/DOKET NUMBER: 37,044

REFERENCE/DOKET NUMBER: 39,044

REFERENCE/REFERENCE/DOKET NUMBER: 39,044

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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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STREET: 1 DNA Way
CITY: South San Fra
STATE: California
COUNTRY: USA
ZIP: 94080
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                                DELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
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                                                                                                                         YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR
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88.7%; Pred. No. 3.7e-157;
tive 21; Mismatches 26;
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US-08-887-352B-16
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Best Local Similarity
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 16:
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TITLE OF INVENTION: Img
TITLE OF INVENTION: Img
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/887,352B
FILING DATE: 03-Jul-1997
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
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CURRENT APPLICATION DATA:
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CITY: South San Francisco
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Amino Acid
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                  YNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSR 357
                                                                                    GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
                                                                                                                                                           SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL 237
                                                                                                                                       SSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELL
                                                                                                                                                                                                        SSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQ 179
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                                                                 GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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1 DNA Way
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Improved Anti-IgE Antibodies and Method of
Improving Polypeptides
26
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TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 65
SEQUENCE CHARACTERISTICS:
LENGTH: 451 amino acids
TYPE: Amino Acid
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Best Local :
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Patent No. 60374
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APPLICANT: Jardieu, Paula M.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/466
FILING DATE: 06-Jun-1995
APPLICATION NUMBER: 08/405
FILING DATE: 15-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/185
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PILING DATE: 07-MAY-: PRIOR APPLICATION DATA: APPLICATION NUMBER: 0
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,0
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CURRENT APPLICATION DATA:
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                 60 YSDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVK--LGTYYFDSWGQGTTLTV 117
                                                                                                            1 EVILVESGGDFVKPGGSLKVSCAASGFAF-SHYAMSWVRQTPAKRLEWVAYISSGGSGTY 59
                                                                                                                                                                        Similarity
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                                                                     EVQLVESGGGLVQPGGSLRLSCAVSGYSITSGYSWNWIRQAPGKGLEWVASITYDGS-TN
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                                                                                                                                                                  69.1%; Score 2122; DB 3; 88.7%; Pred. No. 3.7e-157;
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Sequence 14, Application US/09109207C

Patent No. 6172213

GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
ITILE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptide
FILE REFERENCE: p1123R1
CURRENT FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US/09/109,207C
PRIOR FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 14
LENGTH: 451
TYPE: PAT
ORGANISM: Artificial
FEATURE:
NAME/KEY: Artificial
LOCATION: 1-451
COCHER INFORMATION: Heavy chain sequence derived from MAE11
US-09-109-207C-14
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GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQ
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88.7%;
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Pred. No. 3.7e
21; Mismatches
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3.7e-157;
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Sequence 16, Application US/09109207C
Patent No. 6172213
GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Prest
TITLE OF INVENTION: Improved Anti-IgE Antibo
FILE REFERENCE: P1123R1
CURRENT APPLICATION NUMBER: US/09/109,207C
CURRENT FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/051,554
PRIOR FILING DATE: 1997-07-03
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; LCCATION: 1-451
; OTHER INFORMATION: Heavy chain sequence derived from MAB11
US-09-109-207C-16
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US-09-109-207C-16
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SEQ ID NO 16
; LENGTH: 451
; TYPE: PRT
; ORGANISM: Artificial
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Improved Anti-IgE Antibodies
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; Pred. No. 3.7e-157;
21; Mismatches 26;
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Antibodies and Method of Improving Polypeptide
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Sequence 14, Application US/09296005

Sequence 14, Application US/09296005

Patent No. 6290957

GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptides FILE REFERENCE: P1123C1r

CURRENT APPLICATION NUMBER: US/09/296,005

CURRENT FILING DATE: 1999-04-21

EARLIER APPLICATION NUMBER: US 08/887,352

EARLIER APPLICATION UMBER: US 08/887,352

NUMBER OF SEQ ID NO 14

LENGTH: 451

TYPE: PRT
ORGANISM: Artificial
FEATURE:
NAMES/KEY: Artificial
FEATURE:
NAMES/KEY: Artificial
LOCATION: 1-451

LOCATION: 1-451

LOCATION: 1-451

LOCATION: Heavy chain sequence derived from MAEI1

US-09-296-005-14
Search completed: December 23, 2004, 19:08:03 Job time: 39.4376 secs
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US-09-296-005-14
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                                                                                     RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
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1: /cgn2_6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO6_PUBCOMB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

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Copyright (c) 1993 - 2004 Compugen Ltd.
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/ Cgn2_6/ptodata/2/pubpaa/US09E PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US09 NEW PUB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10E PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10E PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10E PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10E PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10D PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10D PUBCOMB.pep: *
/ Cgn2_6/ptodata/2/pubpaa/US10E PUB.pep: *
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US-10-310-719-32
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US-10-310-719-35
US-10-412-406-32
US-10-679-620-64
US-10-679-620-62
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    Sequence 41, Appl
Sequence 32, Appl
Sequence 6, Appl
Sequence 37, Appl
Sequence 37, Appl
Sequence 32, Appl
Sequence 32, Appl
Sequence 64, Appl
Sequence 64, Appli
Sequence 64, Appli
Sequence 6, Appli
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ALIGNMENTS

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Sequence 41. Application US/10138727A

Publication No. US20030157054A1

GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Susan
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/10/138,727A
CURRENT APPLICATION NUMBER: US/10/138,727A
CURRENT PILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: US/0/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.0

SEQ ID NO 41
                                                                                                                                                                                                                                                                                        ; LENGTH: 579
; TYPE: PRT
; ORGANISM: Artificial sequence
; PEATURE:
; OTHER INFORMATION: heavy chain-IL2
US-10-138-727A-41
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                                           61 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTR-VKLGTYYFDSWGQGTTLTVSS 119
61 ADDFKGRFAFSLETSTSTAFLQINNLRSEDTATYFCVRFISKGDY----WGQGTSVTVSS
                                                                                                                          QIQLVQSGAEVKKPGETVKISCKASGYTFTNYGMNWVKQTPGKGLKWMGWINTYTGEPTY 60
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les 31; Indels
                                                                                                                                                                                                                                          DB 14; Length 579;
                                                                                                                                                                                            5; Gaps
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CURRENT APPLICATION NUMBER: US/10/310,719
CURRENT FILING DATE: 2002-12-04
PRIOR APPLICATION NUMBER: 60/337,113
PRIOR FILING DATE: 2001-12-04
PRIOR PRILING DATE: 2001-12-04
PRIOR FILING DATE: 2002-04-12
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.1
SEQ ID NO 32
LENGTH: 579
TYPE: PRT
ORGANISM: Artificial Sequence
PRINTER: Artificial Sequence
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publication No. US20030166163A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Immunocytokines With Modulated
FILE REFERENCE: LEX-020
                                                                                                                                                                  Query Match
Best Local Similarity
Matches 515; Conserv
                                                                                                                                                                                                                           OTHER INFORMATION: dI-KS-ala IL2 (D20T) heavy chain fused to IL-2 variant-10-310-719-32
                                                                                                                                                                                                                                                      FEATURE:
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           ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                         VIVLELKGSETTFMCEYADETATIVEFLNRWITFCQSIISTLT 582
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                                                    ADDFKGRFTITAETSTSTLYLQLNNLRSEDTATYFCVRFISKGDY---
                                                                              SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTR-VKLGTYYFDSWGQGTTLTVSS
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ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
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                                                                                                                                                                     Conservative
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                                                                                                                                                                ; Score 2738.5; DB 14;
; Pred. No. 3.3e-166;
30; Mismatches 33; I
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FILE REFERENCE: LEX-023
CURRENT APPLICATION NUMBER: US/10/737,208A
CURRENT FILING DATE: 2003-12-16
PRIOR APPLICATION NUMBER: US 60/433,945
PRIOR FILING DATE: 2002-12-17
NUMBER OF SEQ ID NOS: 6
SOPTWARE: Patentin version 3.1
SEQ ID NO 6
LENGTH: 575
                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Humanized Immunoglobulin Heavy Chain-IL-2
US-10-737-208A-6
S
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US-10-737-208A-6
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                                                                                                                                                                                                                                                                                                                                 Best Local Similarity Matches 517; Conserv
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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APPLICANT: Lo, Kin-Ming
TITLE OF INVENTION: IMMUNOCYTOKINE SEQUENCES
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                                                                                                                                                                   NQKFKGRATLTVDKSTSTAYMHLKSLRSEDTAVYYCVS---GMEY---WGQGTSVTVSSA
                                                                                                                                                                                                       SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSSA 120
SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS
                                                                                                            STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                               STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
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Pred. No. 9.5e
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Sequence 37, Application US/10310719

| Publication No. US20030166163A1
| GENERAL INFORMATION: LESHOR
| APPLICANT: Gillies, Stephen
| TITLE OF INVENTION: Immunocytokines With Modulated Selectivity
| FILE REFERENCE: LEX-020
| CURRENT APPLICATION NUMBER: US/10/310,719
| CURRENT FILING DATE: 2002-12-04
| PRIOR APPLICATION NUMBER: 60/337,113
| PRIOR APPLICATION NUMBER: 60/337,113
| PRIOR APPLICATION NUMBER: 60/337,113
| PRIOR APPLICATION NUMBER: 60/371,966
| PRIOR FILING DATE: 2002-04-12
| NUMBER OF SEQ ID NOS: 37
| SOPTWARE: Patentin version 3.1
| SEQ ID NO 37
| LENGTH: 580
| TYPE: PRI
| CREATING DATE: 1 SCREECE
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US-10-310-719-37
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Best Local S
Matches 494
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nes 494; Conservative
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                                                                                                                                                                                                                                                                           ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS 179
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                                                          PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN 299
                                                                                                                                  GLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVDKRVEPKSCDKTHTCPPCPAPEFLGG
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                           PSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQAQ
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Pred. No. 7.1e-157;
11; Mismatches 54;
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US-10-310-719-35
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Publication No. US20030166163A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Immunocytokines With Modulated
FITE BORDER FOR A PROPERTY OF THE CONTROL OF TH
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Best Local
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PRIOR FILING DATE: 2001-12-04
PRIOR APPLICATION NUMBER: 60/371,966
PRIOR FILING DATE: 2002-04-12
NUMBER OF SEQ ID NOS: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: LEX-020
CURRENT APPLICATION NUMBER: US/10/310,719
CURRENT FILING DATE: 2002-12-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: 580
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRD 358
||:||||||||:|||||||:
QSTFRVVSVLTVVHQDWLNGKEYKCKVSNKGLPAPIEKTISKTKGQPREPQVYTLPPSRE 356
                                                                                                                                     | GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQY 298
                                                                                                                                                                                                                                            SGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLG 238
                                                                                                                                                                                                                                                                                                                          GASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YSDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 119
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                                                                                                         GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQA
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84.1%; Pred. No. 7.8e-155;
tive 31; Mismatches 56;
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US-10-412-406-32
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CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: 60/240,285
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/275,289
PRIOR FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: 60/299,987
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: FCT/US01/32140
PRIOR APPLICATION NUMBER: PCT/US01/32140
PRIOR FILING DATE: 2001-10-12
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 418; Conserv
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SEQ ID NO 32
LENGTH: 663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SALDHANA, JOSE W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINALOOCN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 33
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APPLICANT: GARBER, EL
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STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
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                                                                                                       GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG
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                                            PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN
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                                                                                                                                                                                                                                                                                                                                                         72.2%;
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TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R AN FILE REFERENCE: BINALOCCN
CURRENT APPLICATION NUMBER: US/10/412,406
CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: 60/240,285
PRIOR FILING DATE: 2000-10-13
PRIOR FILING DATE: 2001-03-13
PRIOR FILING DATE: 2001-03-13
PRIOR PILING DATE: 2001-06-21
PRIOR PILING DATE: 2001-01-12
PRIOR PILING DATE: 2001-06-21
PRIOR PILING DATE: 2001-06-21
PRIOR PILING DATE: 2001-06-21
PRIOR PILING DATE: 2001-01-12
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US-10-412-406-33
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; LENGTH: 4852
; TYPE: PRT
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Best Local Similarity 93.1%;
Matches 418; Conservative
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APPLICANT: GARBER, Ellen
APPLICANT: LYNE, Paul
APPLICANT: SALDHANA, JOSE
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QQGNVFSCSVMHEALHNHYTQKSLSLSPG 448
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                                                                                     LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
                                                                                                                                                                  STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE
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Pred. No. 5.4e-132;
9; Mismatches 21;
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RESULT 9
US-10-679-620-62
Sequence 62, Application US/10679620
; Publication No. US20040110930A1
; GENERAL INFORMATION:
; APPLICANT: Large Scale Biology
; APPLICANT: Reinl, Stephen J.
; APPLICANT: Edwards, Patricia C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 34150-004A
CURRENT APPLICATION NUMBER: US/10/679,620
CURRENT FILING DATE: 2003-10-03
PRIOR APPLICATION NUMBER: 60/415,940
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 122
SOFTWARE: Patentin version 3.2
SEQ ID NO 64
LEGITH: 713
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
COMMENT TYPESTANTION. PREFICELY TO SEE INC.
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US-10-679-620-64
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Best Local Similarity
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APPLICANT: Reinl, Stephen J.
APPLICANT: Edwards, Patricia C.
TITLE OF INVENTION: MULTIMERIC PROTEIN ENGINEERING
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                                                                                                                                                                                                    TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                                                                                                                    LPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKL 412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCP 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PDSVKGRFTISRDNDKNALYLQMNSLKSEDTAMYYCARRSEFYYYGNTYYYSAMDYWGQG 376
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                                                                                                                                                                                                                                               LPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKL
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                                                                                                                                                                             TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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Pred. No. 8.2e-133;
0; Mismatches 20;
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                                              APPLICANT: BIOGEN, INC.
TITLE OF INVENTION: ANTHODIES TO VLA-1
FILE REFERENCE: A101 PCT
CURRENT APPLICATION NUMBER: US/10/474,832
CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR APPLICATION NUMBER: 60/303,689
PRIOR PILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
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US-10-474-832-4
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LENGTH: 715
TYPE: PRT
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Best Local
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CURRENT APPLICATION NUMBER: US/10/679,
CURRENT FILING DATE: 2003-10-03
PRIOR APPLICATION NUMBER: 60/415,940
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 122
LENGTH: 447
TYPE: PRT
ORGANISM: Artificial Sequence
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91.7%;
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259 EVDLVESGGDLVKPGGSLKLSCAASGFTFSHYGMSWVRQTPDKRLEWVATIGSRGTYTHY 318
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TVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
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Pred. No. 8.3e-133;
10; Mismatches 20;
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APPLICANT: Adolf, G. et al.
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
FILE REFERENCE: 1/221
CURRENT APPLICATION NUMBER: US/10/150,475A
CURRENT FILING DATE: 2002-05-17
PRIOR APPLICATION NUMBER: US 60/307,451
PRIOR FILING DATE: 2001-07-24
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 6
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                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Humanised; OTHER INFORMATION: Murine Antibody BIWA 4 Heavy Chain SEQ ID NO: US-10-150-475A-6
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US-10-150-475A-6
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Best Local S
Matches 416
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                                                                Matches
                                                                             Query Match
Best Local :
                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                     LENGTH: 444
                                                                                                                                                                           FEATURE:
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EVQLVESGGGLVKPGGSLRLSCAASGFTFSSYDMSWVRQAPGKGLEWVSTISSGGSYTYY
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92.9%;
                                                                           71.7%;
                                                           Score 2200.5;
Pred. No. 4.6e
11; Mismatches
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Pred. No. 4e-132;
9; Mismatches 2
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                                                           00.5; DB 14;
. 4.6e-132;
tches 18; I
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APPLICANT: Holder, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and methods for
TITLE OF INVENTION: Cytotoxic CD44 Antibody Imm
FILE REFERENCE: 1/1414
CURRENT APPLICATION NUMBER: US/10/704,522
CURRENT FILING DATE: 2003-11-07
PRIOR APPLICATION NUMBER: US 60/429,516
PRIOR APPLICATION NUMBER: EP 02024881
PRIOR APPLICATION NUMBER: EP 02024881
PRIOR FILING DATE: 2002-11-08
NUMBER OF SEQ ID NOS: 9
SOFTMARE: Patentin Ver. 2.1
SEQ ID NO 6
LENGTH: 444
TYPE: PRT
CURRENT: Artificial Sequence
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                                                                                                                                                                                                                            Matches 415;
                                                                                                                                                                                                                                             Best
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                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Heavy
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                                                                                                                                                         Similarity
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LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
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No. 4.6e-132;
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Immunoconjugates
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US-10-645-215-6
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CURRENT APPLICATION NUMBER: US/10/645,215
CURRENT FILING DATE: 2003-08-21
PRIOR APPLICATION NUMBER: EP 02 018 686.2
PRIOR FILING DATE: August 21, 2002
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 26, 2002
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Heavy Chain US-10-645-215-6
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Best Local S
Matches 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/10645215
Publication No. US20040126379A1
GENERAL INFORMATION:
APPLICANT: Adolf, Guenther
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and Methods for Treating Cancer using
TITLE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and
TITLE OF INVENTION: Chemotherapeutic Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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                                                                                                                                    LYSLSSVYTVPSSSLGTQTY1CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP 240
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    TYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDEL 360
                                                              SVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNS 300
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                                                                                                                  LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGP
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APPLICANT: Bolhuis, Reinier LH
APPLICANT: Woehl, Thorsten
APPLICANT: Woehl, Thorsten
APPLICANT: Boettger, Volker
TITLE OF INVENTION: Wethod of Producing Recombinant Antibodies
FILE REFERENCE: 2923-552
CURRENT APPLICATION NUMBER: US/10/635,908
CURRENT APPLICATION NUMBER: PCT/EP02/01283
PRIOR APPLICATION NUMBER: FCT/EP02/01283
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/266,853
PRIOR APPLICATION NUMBER: 60/266,853
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin version 3.2
SEQ ID NO 16
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US-10-635-908-16
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US-10-635-908-16
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Similarity 92.2%;
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                                                            TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ 420
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10. US20040219633A1
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Pred. No. 7.8e-132;
(4; Mismatches 21;
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RESULT 15
US-10-635-908-18
; Sequence 18, Application US/10635908
; Publication No. US20040219633A1
; GENERAL INFORMATION:
; APPLICANT: Bolhuis, Reinier LH
; APPLICANT: Woehl, Thorsten
; APPLICANT: Boettger , Volker
; TITLE OF INVENTION: Method of Producing Recombinant Antibodies
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SOFTWARE: Patentin version 3.2
SEQ ID NO 18
LENGTH: 449
TYPE: PRT
ORGANISM: Artificial
FEATURE:
Search completed: December Job time: 133.396 secs
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CURRENT APPLICATION NUMBER: US/10/635,908
CURRENT FILING DATE: 2003-08-07
PRIOR APPLICATION NUMBER: PCT/EP02/01283
PRIOR FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/266,853
PRIOR APPLICATION NUMBER: 60/266,853
PRIOR FILING DATE: 2001-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 71.5%; Score 2197; DB 17; Length 449; Best Local Similarity 92.2%; Pred. No. 7.8e-132; Matches 414; Conservative 14; Mismatches 21; Indels 0
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121 STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPBPVTVSWNSGALTSGVHTFPAVLQSSG 180
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Post-processing: Minimum Match 0%
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Gapop 10.0 , Gapext 0.5
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3071
1 EVTLVESGGDFVKPGGSLKV.....IVEFLNRWITFCQSIISTLT 582
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                                                                                                                                                                                                                                          283416
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23		21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	თ	S	4.	ω	N		 	Result
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398	393	329	324	333	289	326	308	329	328	323	328	234	328	255	328	475	474	472	446	374	469	470	444	327	326	377	377	330	Length	
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G3MSM	G1MSM	G3MSC	GIMS	PS0018	G3HUWI	PS0017	C30554	G2GP	147161	GHRB	I47158	PT0207	147160	S31866	147159	S01321	G2MS11	S31459	S40295	869339	S37483	S22080	PC4436	G4HU	G2HU	A60764	A23511	GHHU	ID	
3 chain	gamma-1	3 chain	gamma-1 chain	gamma-2b	gamma-3 heavy	gamma-	heavy chain C		gamma 3 chain	gamma chain C	gamma		gamma 2b	gamma-1 c	gamma 2a	gamma-2b	Ig gamma-2b chain	gamma-1 c	gamma-2a chain		gamma-	5	noclonal antibo	gamma-4	gamma-2 chain	gamma-3 chain	gamma-3	Ig gamma-1 chain C	ption	

A;Residues: 136-154, (0',156-165,'Q',167-176,'Q',178-194,'N',196-197,'D',199-238,'E',240,A;Note: this sequence has the Glm(non-1) markers, 239-Glu and 241-Met R;Ponsting1, H; Hilechmann, N. Hoppe-Seyler's Z. Physiol. Chem. 357, 1571-1604, 1976 A;Title: Die Primaerstruktur eines monoklonalen IgG1-Immunglobulins (Myelomprotein Nie),	,'Q',167- 1m(non-1) . 357, 15	6-165 the G n, N. Chem ur ei	136-154, Q', 156-165, sequence has the GI. H.; Hilschmann, N. 's Z. Physiol. Chem. Primaerstruktur ein	A;Residues: 136-154 A;Note: this sequen R;Ponstingl, H.; Hi Hoppe-Seyler's Z. P A;Title: Die Primae	A;Residues: A;Note: this A;Notestingl; R;Ponstingl; Hoppe-Seyler A;Title: Die
R;Rutishauser, Ü.; Cunningham, B.A.; Bennett, C.; Konigsberg, W.H.; Edelman, G.M. Biochemistry 9, 3171-3181, 1970 A;Title: The covalent structure of a human gammaG-immunoglobulin. VIII. Amino acid A;Reference number: A90564; MUID:71064025; PMID:5530842 A;Contents: Bu A;Accession: Bu A;Accession: Bu A;Accession: Parotein	B.A.; Ber of a hum D:7106402	ham, 1970 cture ; MUI	Cunning 1-3181, nt stru A90564	R;Rutishauser, Ū; Biochemistry 9, 317 B;Tile: The covale A;Title: The covale A;Reference number: Bi A;Contents: Bu A;Accession: A90564 A;Molecule type: px	R;Rutis Biochem A;Tile A;Refer A;Conte A;Acces A;Molec
Biochemistry 9, 3161-3170, 1970 A;Title: The covalent structure of a human gammaG-immunoglobulin. VII. Amino acid A;Reference number: A90563; MUID:71064024; PMID:5489771 A;Contents: myeloma protein Eu A;Accession: B90563 A;Accession: B90563 A;Molecule type: protein A;Residues: 1-96, 'R', '98-135 <cun> A;Note: this sequence has the Glm(3) marker, 97-Arg</cun>	of a hun D:7106402 N> Nm ar	1970 cture ; MUI n Eu 5 <cu< td=""><td>1-3170, nt stru A90563 protei otein ',98-13</td><td>Biochemistry 9, 3161-3170, 1970 A;Title: The covalent structure of a A;Reference number: A90563; MUID:710 A;Contents: myeloma protein Eu A;Accession: B90563 A;Molecule type: protein A;Molecule type: protein A;Mote: this sequence has the Glm(3)</td><td>Biochem A;Title A;Refer A;Reces A;Acces A;Molec A;Note:</td></cu<>	1-3170, nt stru A90563 protei otein ',98-13	Biochemistry 9, 3161-3170, 1970 A;Title: The covalent structure of a A;Reference number: A90563; MUID:710 A;Contents: myeloma protein Eu A;Accession: B90563 A;Molecule type: protein A;Molecule type: protein A;Mote: this sequence has the Glm(3)	Biochem A;Title A;Refer A;Reces A;Acces A;Molec A;Note:
oglobulin gamma genes: implications for evolution of a:83001943; PMID:6811139 U.; Gall, W.E.; Gottlieb, P.D.; Waxdal, M.J.; Edelman,	unoglobul D:8300194	n imm ; MUI ; MUI TAK 17370	of huma \$33887 A 235-330 EMBL:Z Rutish	A;Title: Structure of human immun A;Reference number: S33887; MUID A;Accession: S33887 A;Accession: S33887 A;Molecule type: DNA A;Residues: 88-113;235-330 <tak> A;Cess-references: EMBL:Z17370 R;Cunningham, B.A.; Rutishauser,</tak>	A;Title A;Refer A;Acces A;Molec A;Resid A;Cross R;Cunni
R;Harris, L.J. submitted to the EMBL Data Library, October 1992 A;Reference number: S33904 A;Recession: S36861 A;Accession: S36861 A;Molecule type: DNA A;Residues: 2-330 <har> A;Residues: 2-330 <har> A;Cross-references: EMBL:Z17370 R;Takahashi, N: Ueda, S.; Obata, M.; Nikaido, T.; Nakai, S.; Honjo, T. Cell 29, 671-679, 1982</har></har>	ary, Octo	Libr 17370 Obat	BL Data S33904 A A HAR> EMBL:Z da, S.;	R;Harris, L.J. A;Reference number: S33904 A;Reference number: S33904 A;Recession: S36861 A;Accession: DNA A;Molecule type: DNA A;Molecule type: DNA A;Residues: 2-330 <har> A;Cross-references: EMBL:Z17370 R;Takahashi, N: Ueda, S:; Obata, M:; Nikaido, TCell 29, 671-679, 1982</har>	R; HART18, submitted A; Referen A; Accessi A; Molecul A; Residue A; Cross-r R; Takahas
man immunoglobulin C-gammal gene. 8; PMID:6287432 ::217370 lotypic marker, 97-Lys, and the Glm(1) markers	D:8227423 D:8227423 D:8277423 D:857; EMBI Im(17) al	quenc; MUI; MUI T:P01 the G	A93433 A93433 A ELL> UNIPRO Ce has removed	A;Title: The nucleotide sequence of a human imm A;Reference number: A93433; MUID:82274238; PMID A;Recession: A93433 A;Molecule type: DNA A;Residues: 1-330 <ell> A;Residues: 1-330 <ell> A;Rosidues: 1-330 <els A;Note: this sequence has the Glm(17) allotypic A;Note: Lys-30 is removed after translation</els </ell></ell>	A;Title A;Refer A;Acces A;Molec A;Resid A;Note: A;Note:
Ig gamma-1 chain C region - human G;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 31-Jan-1981 #sequence revision 18-Aug-1982 #text change 09-Jul-2004 C;Date: 31-Jan-1981 #sequence 1533887; B90563; A90564; B91668; A91723; A02146 C;Accession: A9343; S36861; S33887; B90563; A90564; B91668; A91723; A02146 R;Ellison, J.W.; Berson, B.J.; Hood, L.E. Nucleic Acids Res. 10, 4071-4079, 1982	evision 1 3887; B90 Hood, L.I	- hum an) nce r nce r 1; S3	region iens (m #seque ; S3686 rson, B	Ig gamma-1 chain C region - human C;Species: Homo sapiens (man) C;Date: 31-Jan-1981 #sequence rev C;Date: 31-93433; S36861; S338 C;Accession: A93433; S36861; S338 R;Ellison, J.W.; Berson, B.J.; Ho Nucleic Acids Res. 10, 4071-4079;	Ig gamm C;Speci C;Date: C;Acces R;Ellis Nucleic
AL IGNMENTS	ALIC			P	RESULT 1
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A;Reference number: A91668; MUID:77070269; PMID:826475

A;Contents myeloma protein Nie

A;Accession: B91668

A;Molecule type: protein

A;Residues: 1-34, (Y, )46-96, 'K', 98-115,'Q',117-197,'D',199-238,'D',240,'L',242-268,'E',27

A;Molecule type: protein

A;Residues: 1-34, (Y, )46-96, 'K', 98-115,'Q',117-197,'D',199-238,'D',240,'L',242-268,'E',27

A;Molecule type: protein

A;Residues: 1-34, (Y, )46-96, 'K', 98-115,'Q',117-197,'D',199-238,'D',240,'L',242-268,'E',27

A;Note: this sequence has the Glm(1) and Glm(1) markers

A;Contents: myeloma protein KOL; disulfide bonds

A;Title: Die Primaerstruktur des kristallisierbaren monoklonalen Immunglobulins IgG1 KOL

A;Reference number: A91723; MUID:83389131; PMID:8848994

A;Contents: myeloma protein KOL; disulfide bonds

A;Accession: A91723

A;Molecule type: protein

A;Residues: 1-96, R', 98-197,'D',199-238,'E',240,'M',242-266,'D',268-271,'D',273-330 <SCH

A;Residues: 1-96, R',98-197,'D',199-238,'E',240,'M',242-266,'D',268-271,'D',273-330 <SCH

A;Residues: 1-96, R',98-197,'D',
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A;Genetics:
A;Map postation:
14(1; 224/1)
C;Complex:
An immunoglobulin beterotetramer subunit consists of two identical light (kap hain disulfide bonds:
In some cases, such as IgA and IgW, the subunits associate into la C;Cuperfamily:
Immunoglobulin C region; immunoglobulin homology
C;Keywords:
G;Superfamily:
Immunoglobulin homology <IM1>
F;20-85/Domain:
Immunoglobulin homology <IM2>
F;137-206/Domain:
A;Map postation:
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Pred. No. 2.6e-92;
0; Mismatches 0;
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Ig gamma-3 chain C region (allotype G3m(b)) - human C;Species: Homo sapiens (man) C;Date: 28-Dec-1987 #sequence_revision 28-Dec-1987 #text_change C;Accession: A23511 R;Huck, S.; Fort, P.; Crawford, D.H.; Lefranc, M.P.; Lefranc, G. Nucleic Acids Res. 14, 1779-1789, 1986 Nucleic Acids Res. 14, 1779-1789, 1986 A;Title: Sequence of a human immunoglobulin gamma 3 heavy chain (A;Reference number: A23511; MUID:86148507; PMID:3081877 A;Accession: A23511
A;Molecule type: DNA A;Residues: 1-377 <HUC-A,;Cross-references: GB:X03604; GB:M12958; NID:g33070; PIDN:CAA277 C;Genetics:
                                                                                                                                                                   Ig gamma-3 chain C region, form LA C;Species: Homo sapiens (man) C;Date: 14-May-193 #sequence_revi C;Accession: A60764 R;Huck, S.; Lefranc, G.; Lefranc, Immunogenetics 30, 250-257, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Gene: GDB:IGHG3
A;Cross-references: GDB:119339; OMIM:147120
A;Cross-references: GDB:119339; OMIM:147120
A;Map postion: 14q32.33-14q32.33
A;Introns: 98/3; 115/3; 130/3; 145/3; 160/3; 270/3
C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: immunoglobulin homology <IMM>
A;Cross-references: UNIPROT:Q8N4Y9
C;Superfamily: immunoglobulin C re
C;Keywords: immunoglobulin
                                                     A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-377 <HUC>
                                                                                                             Immunogenetics 30, 250-257, 1989
A;Title: A human immunoglobulin IGHG3 allele
A;Reference number: A60764; MUID:90007613; PM
A;Accession: A60764
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ilarity 82.0%;
Conservative 10
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                     region;
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Pred. No. 1.4e
LO; Mismatches
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;20-85/Domain:

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A; Molecule type: protein

A; Residues: 1-19,'Q', 21-57,'Z',59,'A',61-193,'D',195-325 <WANNAR, Residues: 1-19,'Q',21-57,'Z',59,'A',61-193,'D',195-325 <WANNAR, Residues: Trp-156 is at or near the complement-binding site

R; Connell, G.E.; Parr, D.M.; Hofmann, T.

Can. J: Biochem. 57, 758-767, 1979

A; Title: The amino acid sequences of the three heavy chain cone

A; Reference number: A90752; MUID:80001357; PMID:113060

A; Contents: myeloma protein Zie

A; Molecule type: protein

A; Residues: 1-24,'E',26-57,'EV',60-85;132-171,'ZZZ',175,'B',177

A; Note: this sequence has since been revised

R; Hofmann, T.; Parr, D.M.

Mol. Immunol. 16, 923-925, 1979

Mol. Immunol. 16, 923-925, 1979
   A;Title: A note on the amino acid sequence A;Reference number: A93132; MUID:80114419; A;Contents: Zie
                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:P01859; GB:V00554; GB:J00230; NI
A;Note: Lys-326 is probably removed posttranslationally
R;Wang, A.C.; Tung, E.; Eudenberg, H.H.
J. Immunol. 125, 1048-1054, 1980
A;Title: The primary structure of a human IgG2 heavy chain:
A;Reference number: A92809; MUID:81007873; PMID:6774012
A;Contents: myeloma protein Til
A;Accession: A92809
A;Accession: A92809
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C;Date: 30-Apr-1981 #sequence revision 13-Jun-1983 #text_change 09-
C;Accession: A93906, A92809; A90752; A93132; A02148
R;Ellison, J.; Hood, L.
Proc. Natl. Acad. Sci. U.S.A. 79, 1984-1988, 1982
A;Title: Linkage and sequence homology of two human immunoglobulin A;Reference number: A93906; MUID:82197621; PMID:8804948
A;Accession: A93906
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A;Residues: 1-326 <ELL>
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Pred. No. 1.9e
LO; Mismatches
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PMID:118920
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.9e-84;
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                                          of human
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                                        immunoglobulin
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Ig gamma-4 chain C region - human c;Species: Homo sapiens (man) C;Date: 02-Apr-1982 #sequence revision C;Accession: A90933; A90249; A02150 R;Ellison, J.; Buxbaum, J.; Hood, L. DNA 1, 11-18, 1981 DNA 1, 11-18, 1981
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R;Milstein, C.; Frangione, B.
Biochem. J. 121, 217-225, 1971
A;Title: Disulphide bridges of the heavy chain of A;Reference number: A90253; MUID:72033500; PMID:4: A;Contents: annotation; myeloma protein Sa, disul: R;Frangione, B.; Milstein, C.; Pink, J.R.L.
Nature 221, 145-148, 1969
A;Title: Structural studies of immunoglobulin G.
A;Reference number: A93157; MUID:69064124; PMID:5.
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A; Molecule type: protein
A; Residues: 238-275 <HOF>
R; Hofmann, T.; Parr, D.M.
submitted to the Atlas, Maj
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C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (ka C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (ka hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into 1 C;Superfamily: immunoglobulin C region; immunoglobulin homology C;Keywords: duplication; glycoprotein; heterotetramer; immunoglobulin F;20-85/Domain: immunoglobulin homology <IM1>
F;133-202/Domain: immunoglobulin homology <IM2>
F;239-306/Domain: immunoglobulin homology <IM3>
F;239-306/Domain: immunoglobulin homology <IM3>
F;147Disulfide bonds: interchain (to light chain) #status experimental F;27-83,140-200,246-304/Disulfide bonds: #status experimental F;102,103,106,109/Disulfide bonds: interchain (to heavy chain) #status experimental F;176/Binding site: carbohydrate (Asn) (covalent) #status predicted
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01; Conservative
                      QQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                      LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW 419
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                                                                                          MTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDKSRW
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Pred. No. 3.7e-83;
2; Mismatches 13
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that shown in having
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PMID:4940472
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PMID:5782707
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human

immunoglobulin

C-gamma4

02-Apr-1982

#text_change

09-Jul-2004

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A;Gene: GDB:IGHG4
A;Gene: GDB:IGHG4
A;Cross-references: GDB:119340; OMIM:147130
A;Map position: 1492.33-14932.33
A;Introns: 99/1; 11/1; 221/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: duplication; glycoprocein; heterotetramer; immunoglobulin
F;20-85/Domain: immunoglobulin homology <IM1>
F;39-110/Region: hinge
F;134-203/Domain: immunoglobulin homology <IM2>
F;240-307/Domain: immunoglobulin homology <IM3>
F;471sulfide bonds: interchain (to light chain) #status experimental
F;27-83,141-201,247-305/Disulfide bonds: #status predicted
F;106,109/Disulfide bonds: interchain (to heavy chain) #status experimental
F;177/Binding site: carbohydrate (Asn) (covalent) #status predicted
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A;Accession: A90933
A;Molecule type: DNA
A;Residues: 1-327 <ELL>
A;Residues: 1-327 <ELL>
A;Cross references: UNIRROT:P01861
A;Cross references: UNIRROT:P01861
A;Note: the sequence was determined from the (R;Pink, J.R.L.; Buttery, S.H.; De Vries, G.M., Biochem. J. 117, 33-47, 1970
A;Title: Human immunoglobulin sublclasses. Para,Reference number: A90249; MUID:70207560; PM.
A;Accession: A90249
A;Molecule type: protein
A;Residues: 1-30;81-326 <PIN>
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                               RESULT
PC4436
DDDDBBCCCC
                                                                                                                                                                                                                                                               monoclonal antibody 13-1 heavy chain - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 04-Feb-1998 #sequence_revision 13-Mar-1998 #text_change
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Matches 300
                            Reference number: JC5810; MUID:98063277; PMID:9398605;Accession: PC4436;Molecule type: protein
                                                                                                                              Accession: PC4436
Accession: PC4436
Accession: Kato, K.; Torizawa, T.; Dohmae, N.; Yamaguciochem. Biophys. Res. Commun. 240, 566-572, 1997
Title: Structural characterization of mouse monoclonal
Residues:
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Pred. No. 2.1e
12; Mismatches
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G.M.; Milstein, C.
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1:1e-82;
nes 15;
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A;Cross-references: EMBL:X16701
A;Note: the sequence was determined from the C;Genetics:
A;Gene: Ig CH gamma-1
A;Introns: 98/1; 111/1; 221/1
C;Superfamily: immunoglobulin C region; immu C;Superfamily: immunoglobulin homology <I
                                                                                                                                                                                                         A;Molecule type: mRNA
A;Residues: 1-470 <SAN4
A;Residues: 1-470 <SAN4
A;Residues: 1-470 <SAN4
A;Residues: EMBL:X62916; NID:9439; PIDN:CAA44699.1;
A;Symons, D.B.A.; Clarkson, C.A.; Beale, D.
Mol. Immunol. 26, 841-850, 1989
Mol. Immunol. 26, 841-850, 1989
A;Title: Structure of bovine immunoglobulin constant region
A;Reference number: S06610; MUID:90097956; PMID:2513487
                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 7

$22080

Heavy chain precursor (B/MT.4A.17.H5.A5) - bovir Ig heavy chain precursor (B/MT.4A.17.H5.A5) - bovir Ig heavy chain precursor (B/MT.4A.17.H5.A5) - bovir Ig heavy chain precursor (chain C region (clone C,Species: Bos primigenius taurus (cattle) C,Date: 06-Jan-1995 #sequence revision 06-Jan-1995 C,Accession: $22080; $06610; $31303

R,Sanders, P.G.
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                                                                                                                                                A; Molecule type: DNA
A; Residues: 142-470 <SYM>
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Pred. No. 1.4
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                     immunoglobulin;
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submitted to the EMBL Data Library, February 1993
A;Reference number: S37483
A;Accession: S37483
A;Statue: preliminary
A;Molecule type: mRNA
A;Residues: 1-469 < DUC>
A;Cross-references: EMBL:X70423; NID:g406252; PIDN:CAA49868.1;
C;Superfamily: immunoglobulin
C;Reywords: immunoglobulin
C;Reywords: immunoglobulin
F;276-345/Domain: immunoglobulin homology <IMM>
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$37483

Ig gamma-2a chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision
C;Accession: $37483
R;Ducancel, F.F.D.
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Best Local Similarity
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                                   ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS 179
                                                                                                                                                        DKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QSSGLYSLSSMVTVPGSTSG-QTFTCNVAHPASSTKVDKAVDP-TC-KPSPCDCCPPPEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NPALKSRLSITKENSKSQVSLSVSSVTPEDTATYYCARSTYGEVGDGAIADAWGQGLLVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
              AKTTAPSVYPLAPVCGDTTGSSVTLGCLVKGYFPEPVTLTWNSGSLSSGVHTFPAVLQSD
                                                                                                            SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGT-YYFDSWGQGTTLTVSS
                                                                                                                                                                                            EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PGGPSVFIFPPKPKDTLTISGTPEVTCVVVDVGHDDPEVKFSWFVDDVEVNTATTKPREE
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                                                                                     NENFKGKATLTVDTSSSTAYMOLSSLTSEDTAVYFCARAMGATATLLDYWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DRNSWQEGDTYTCVVMHEALHNHYTQKSTSKSAGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTY----YFDSWGQGTTLT 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QVQLRESGPSLVKPSQTLSLTCTVSGFSLSSYALTWVRQAPGKALEWVGGITSGGT-TYY
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                                                                                                                                                                                                                             Score 1430.5; DB Pred. No. 3.2e-73; 71; Mismatches 110
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Pred. No. 3.3e-75;
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                                                                                                                                                                                                                                                                     469;
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$69339
Ig heavy chain V region precursor - human
C;Species: Homo sapiens (man)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change |
C;Accession: $69339; $72664
C;Accession: $69339; $72664
R;Khamlichi, A.A.; Aucouturier, P.; Preud'homme, J.L.; Cogne, M.
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A;Title: Structure of abnormal heavy chains in human heavy-chain-deposition
A;Reference number: S69339; MUID:95262687; PMID:7744049
A;Accession: S69339
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A; Residues: 1-140,'C',142-374 < KH2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Reference number: A; Accession: S72664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Khamlichi, A.A.
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A; Residues: 1-374 < KHA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loc
Matches
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;Superfamily: immunoglobulin C region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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hes 281;
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EQYNSTYRVSVLTVLHQDWLNGKEYKCKVSNKALFAFIEKTISKAKGQPREFQVYTLFF
                  EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
                                                                      LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
                                                                                       LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWXVDGVEVHNAKTKPRE
                                                                                                                                                                  LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPE
                                                                                                                                                                                                                                                TVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
                                                                                                                                                                                                                                                                                    RYSPSLRTRLTITKDTSKNQVVLTMTNVDPADTATYYCGYSVEGYGQGYRFHSWGQGTLV
                                                                                                                                                                                                                                                                                                                 YYSDSVKGRFTISRDNAKOTLYLOMRSLRSEDSAMYFCTRVKLG----TYYFDSWGQGTTL
                                                                                                                                                                                                                                                                                                                                                          QITLKESGPTLVKPTQTLTLTCTFSGFSLSKSGVGVGWIRQPPGQALEWLALI-FWDDDK 78
                                                                                                                                                                                                                                                                                                                                                                                            EVTLVESGGDFVKPGGSLKVSCAASGFAFSH--YAMSWVRQTPAKRLEWVAYISSGGSGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                46.3%; Score 1423; DB 2; llarity 61.9%; Pred. No. 6.4e-73; Conservative 23; Mismatches 46;
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A; Map position: 12
C; Superfamily: immunoglobulin C region; immunoglobulin homology.
C; Keywords: disulfide bond; glycoprotein; immunoglobulin; pyrog.
F; 1-446/Product: Ig gamma-2a chain #status experimental 
F; 1-117/Domain: V-D-J region 
F; 1-117/Domain: V-D-J region 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;22-96,144-199,261-321,367-425/Disulfide bonds: #status predicted F;132/Disulfide bonds: interchain (to light chain) #status predicted F;224,227,229/Disulfide bonds: interchain #status predicted F;224,227,229/Disulfide bonds: interchain #status predicted F;297/Binding site: carbohydrate (Asn) (covalent) #status experimental
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F;118-214/Domain: C1 region <CH1>
F;215-230/Region: hinge
F;231-340/Domain: C2 region <CH2>
F;341-446/Domain: C3 region <CH3>
F;360-427/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Reference number: S40295
A; Accession: S40295
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A;Description: Primary structure of the murine monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ś
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Best Local
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                                                                                           NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRD
                                                                                                                                                              GPSVFLFPPKFKDTLMISRTPEVTCVVVDVSHEDFEVKFNWYVDGVEVHNAKTKFREEQY
                                                                                                                                                                                                                                          LYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPP--CPAPELLG
                                                                                                                                                                                                                                                                                       KTTAPSVYPLAPVCGDTTGSSVTLGCLVKGYFPEPVTLTWNSGSLSSGVHTFPAVLQSD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMTKKQVTLTCMVTDFMPEDIYVEWTNNGKTELNYKNTEPVLDSDGSYFMYSKLRVEKKN
                GPSVFIFPPKIKDVLMISLSPMVTCVVVDVSEDDPDVQISWFVNNVEVLTAQTQTHREDY
                                                                                                                                                                                                                    LYTLSSSVTVTSSTWPSQSITCNVAHPASSTKVDKKIEPRG-PTIKPCPPCKCPAPNLLG
                                                                                                                                                                                                                                                                                                                          STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSG
                                                                                                                                                                                                                                                                                                                                                               NEKFKGKATLTVDTSSSTAYMQLSSLTSEDSAVYFCARG--GKFAMDYWGQGTSVTVSSA
                                                                                                                                                                                                                                                                                                                                                                                                                                    QIQLQQSGPELVRPGASVKISCKASGYTFTDYYIHWVKQRPGEGLEWIGWIYPGSGNTKY
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                                                                      NSTLRVVSALPIQHQDWMSGKEFKCKVNNKDLPAPIERTISKPKGSVRAPQVYVLPPPEE
                                                                                                                                                                                                                                                                                                                                                                                               SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSSA
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Pred. No. 1e-72;
Pred. 109;
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C;Species: Ovis orientalis aries, Ovis am
C;Date: 13-Jan-1995 #sequence_revision 13
C;Accession: S31459
R;Patri, S:; Nau, F.
submitted to the EMBL Data Library, Decem
A;Reference number: S31459
A;Accession: S31459
A;Accession: S31459
A;Status: preliminary
A;Malocill tro. EMBL
                                              Ig gamma-2b chain - mouse
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 31-Mar-1980 #sequence revision 01-Dec-2000 #text change C;Accession: S25057; A02157; A26235; A26232; A26233; A53598
R;Fischer, R; Voss, A.; Niersbach, M.; Munziker, W.; Hirsch, H. submitted to the EMBL Data Library, July 1992
A;Description: Production of a Tobacco mosaic virus (TMV) inacti
                                                                                                                                                                              RESULT 12
G2MS11
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A;Residues: 1-472 <PAT>
A;Cross-references: EMBL:X69797
A;Cross-references: EMBL:X69797
C;Superfamily: immunoglobulin
C;Keywords: immunoglobulin
F;277-346/Domain: immunoglobulii
                A; Reference number: S25057
A; Accession: S25057
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preliminary
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Pred. No. 3.5e-
66; Mismatches
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13-Jan-1995
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3.5e-72;
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1 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSG---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QGTTLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVH
                      YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                             YVLAPPQEELSKSTLSVTCLVTGFYPDYIAVEWQKNGQPESEDKYGTTTSQLDADGSYFL
                                                                                                                      YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQP--ENNYKTTPPVLDSDGSFFL
                                                                                                                                                                                           TKPREEQFNSTFRVVSALPIQHQDWTGGKEFKCKVHNEALPAPIVRTISRTKGQAREPQV
                                                                                                                                                                                                                 TKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV
                                                                                                                                                                                                                                                                                    CPPPELPGGPSVFIFPPKPKDTLTISGTPEVTCVVVDVGQDDPEVQFSWFVDNVEVRTAR
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                                                                                                                                                                                                                                                                                                                                                                                                                              TFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPP
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YSRLRVDKNSWQEGDTYACVVMHEALHNHYTQKSISKPPGK
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inactivating H.J.;

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09-Jul-2004

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A;Title: Mouse immunoglobulin allotypes: post-duplication divergence of gamma2a and gamma A;Reference number: A26233; MUID:82173203; PMID:6803173
A;Contents: b allele
A;Accession: A26233
A;Contents: DNA
A;Contents: A3598; MUID:94216359; PMID:7512967
A;Contents: Pype: protein
A;Reference number: A3598; MUID:94216359; PMID:7512967
A;Contents: Pype: Pype: P
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A;Contents: a allele
A;Accession: A02157
A;Contents: a allele
A;Accession: A02157
A;Contents: a allele
A;Accession: A02157
A;Molecule type: DNA
A;Residues: 138-161,'L',163-189,'FP',193-474 <YAM>
A;Residues: 138-161,'L',163-189,'FP',193-474 <YAM>
A;Residues: 138-161,'L',163-189,'FP',193-474 <YAM>
A;Conserverences: GB:J00461
A;Conserverences: GB:J00461
A;Note: the sequence was determined from the germline gene
R;Tucker, P.W.; Marcu, K.B.; Slightom, J.L.; Blattner, F.R.
Science 206, 1299-1303, 1979
A;Title: Structure of the constant and 3' untranslated regions of the murine gamma2b he
A;Reference number: A26235; MUID:80081501; PMID:117548
A;Contents: MPC 11
A;Accession: A26235
A;Molecule type: mRNA
A;Residues: 138-172,'P',174-189,'FP',193-376,'T',378-474 <TU1>
A;Note: Lys-474 is probably removed posttranslationally
R;Tucker, P.W.; Marcu, K.B.; Newell, N.; Richards, J.; Blattner, F.R.
Science 206, 1303-1306, 1979
A;Title: Sequence of the cloned gene for the constant region of murine gamma2b immunogl
A;Residues: 138-172,'P',174-189,'FP',193-376,'T',378-474 <TU2>
A;Molecule type: DNA
A;Residues: 138-172,'P',174-189,'FP',193-376,'T',378-474 <TU2>
Noture 296, 761-763, 1982
A;Title: Marcu, Midd: Midd
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A;Roslidues: 1-474 <FIS>
A;Cross-references: UNIPROT:P01866; EMBL:X67210; NID:g54826; PIDN:CAA47649.1;
A;Cross-references: UNIPROT:P01866; EMBL:X67210; NID:g54826; PIDN:CAA47649.1;
R;Yamawaki-Kataoka, Y.; Kataoka, T.; Takahashi, N.; Obata, M.; Honjo, T.
Nature 283, 786-789, 1980
Nature 283, 786-789, 1980
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NEKFKGKATLTSDKSSNTAYMELSSLTSEDSAVYYCAR----DYDYDWFAYWGQGTLVTV
                                                                                                                                                                                                                                        EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                          SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTY---YFDSWGQGTTLTV
                                                                                                                                                     EVQLQQSGPELVNPGASVKMSCKASGYTFITYVMHWVKQKPGQGLEWIGYINPNKDGTKF
                                                                                                                                                                                                                                                                                                                                                    44.48;
                                                                                                                                                                                                                                                                                                                                                        Score 1365; DB 1;
Pred. No. 1.6e-69;
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F;1-19/Domain: signal sequence #status predicted <SIG>F;20-475/Product: Ig gamma-2b chain #status predicted F;159-223/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:X13188; NID:g51780; PIDN:CAA31580.1; PID:g51781
A;Note: this sequence was determined from the differentiated gene
C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-475 < DE1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Title: Expression in non-1
A; Reference number: S01320;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;de Waele, P.; Feys, V.; van de Voorde,
Eur. J. Biochem. 176, 287-295, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig gamma-2b chain precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 30-Sep_1989 #sequence_revision 30-Sep-1989 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: S01321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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PREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREEQVYT
                                                                                                              APELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTK
                                                                                                                                                                                                             GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKT-HTCPP-----CP
                                                                                                                                                                                                                                                             AKTTPPSVYPLAPGCGDTTGSSVTLGCLVKGYFPESVTVTWNSGSLSSSVHTFPALLQ-S
                                                                                                                                                                                                                                                                                                                                                                                               SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVK-LGTYYFDSWGQGTTLTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS
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                                                                       APNIEGGPSVFIFPPNIKDVLMISLTPKVTCVVVDVSEDDPDVQISWFVNNVEVLTAQTQ
                                                                                                                                                                  GLYTMSSSVTVPSSTWPSQTVTCSVAHPASSTTVDKKLEPSGPTSTINPCPPCKECHKCP
                                                                                                                                                                                                                                                                                                        ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS
                                                                                                                                                                                                                                                                                                                                                    NEKFKGKATLTVDKSSSTAYLHLSSLTSEDSAVYFCAGPRQVGLLPFGYWGQGTLVTASA
                                                                                                                                                                                                                                                                                                                                                                                                                                               QVQLQQSGAELARPGASVKLSCKASGYTLTSYGISWVKQRTGQGLEWIGEIYPGSGNSYF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KLNMKTSKWEKTDSFSCNVRHEGLKNYYYLKKTISRSPGK
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55.4%; Pre/
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Pred. No. 2.5e
75; Mismatches
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RESULT 14

147159

Ig gamma 2a chain constant region - pig (fragment)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_chan;
C;Accession: I47159
R;Kacskovics, I; Sun, J; Butler, J.E.
J. Immunol. 153, 3565-3573, 1994
A;Title: Five putative subclasses of swine IgG identified from A;Reference number: I47158; MUID:95015845; PMID:7930579
A;Reference number: I47158; MUID:95015845; PMID:7930579
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-328 <KAC'>
A;Cross-references: EMBL:U03779; NID:9433123; PIDN:AAA52217.1
C;Genetics:
A;Gene: IGG2a
C;Superfamily: immunoglobulin C region; immunoglobulin homology <IMM>
Ig gamma-1 chain C region - synthetic
C;Species: synthetic
A;Note: Homo sapiens (man) gene engineered and expre
C;Date: 06-Jan-195 #sequence_revision 17-Mar-1997 #
C;Accession: S31866
R;Filpula, D.
Submitted to the EMBL Data Library, February 1993
A;Reference number: S31866
A;Accession: S31866
A;Accession: S31866
A;Molecule type: mRNA
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                                                                                                                                                                                                                                        RESULT
                                                                                                                   g gamma-1 chain C region - synthetic
;Species: synthetic
;Note: Homo sapiens (man) gene engineered and expressed in Escherichia coli
;Date: 06.-Jan-1995 #sequence_revision 17-Mar-1997 #text_change 19-May-2000
;Accession: S31866
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;Superfamily: immunoglobulin C region; immunoglobulin
;Superfamily: immunoglobulin homology <IMM>
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Best Local Similarity
Matches 231; Conserv
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NMKTSKWEKTDSFSCNVRHEGLKNYYLKKTISRSPGK 475
                                                                                                                                                                                                                                                                                                                   RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK 449
                                                                                                                                                                                                                                                                                                                                                                                           LTKNQVSLTCLVKGFYPSDIAVEWESNGQ--PENNYKTTPPVLDSDGSFFLYSKLTVDKS
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                                                                                                                                                                                                                                                                                               SWQGGGIFQCAVMHEALHNHYTQKSISKTPGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APKTAPSVYPLAPCSRDTSGPNVALGCLASSYFPEPVTVTWNSGALSSGVHTFPSVLQPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LSPPPEQLSRKDVSLTCLAVGFSPEDISVEWTSNGHTEENYKDTAPVLDSDGSYFIYSKL
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Pred. No. 5.3e-64;
                                                             protein-protein
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                                                               interactions
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A;Residues: 1-255 <FIL>
A;Cross-references: EMBL:X70421; NID:g33068; PIDN:CAA49866.1; PID:C;Keywords: immunoglobulin
E;1-22/Region: Bscherichia coli outer membrane protein A precursor
E;23-255/Region: human Ig gamma-1 chain C region
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Best Local Similarity
Matches 233; Conserv
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                                                                                    331 PAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPE
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197
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                NNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                     TVAQADVESKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSH
                                                                                                                                                                                                                              TKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSH
                                                                                                                                                      EDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKAL
NNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                PAPIEKTISKAKGOPREPQVYTLPPSRDELTKNOVSLTCLVKGFYPSDIAVEWESNGOPE
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97.5%;
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Pred. No. 5.8e-64;
0; Mismatches 6
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Search completed: December 23, 2004, 19:06:08
Job time : 33.449 secs

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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2004 Compugen Ltd
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Q65DAH

AAH18747

Q6PI81

AAH41037

Q6M26

CAE45972

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Q6MW7

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AAH0670

ACB45781

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SHYAMSWVRQTPAKRLEWVAYISSGGSGTYY 60 : : ::: SSYSMNWVRQAPGKGLEWVSSMSSSSSYIYY 79	; Length 471; Indels 3; Gaps 2;	databases. E CRC64;	woorigues s., sanchez A., o Y., Bouffard G.G., Dickson M.C., ars R.M., Butterfield Y.S., Schnerch A., Schein J.E., an 15,000 full-length human 3(2002).	.G., Schuler G.D., Schuler G.D., Hat N.K., Heigh F., Hong L., Scheetz T. F., Scheetz T. D., Mullahy C., Mullahy S. D., Mullahy S. L.J., Hulyk S. Dbs R.A., L.J., Sanchez	te) ta; Euteleostomi; dae; Homo.	Bac85350 homo sapi Aah14667 homo sapi Aah19337 homo sapi Bac85175 homo sapi Bac85171 homo sapien Cae45773 homo sapien Q727p5 homo sapien Aah62336 homo sapi Bac86255 homo sapi Bac86255 homo sapi G6n095 homo sapien Cae45775 homo sapien

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RETISUE=Primary B-Cells;

REDLINE=22388257; PubMed=12477932;

REDLINE=22388257; PubMed=12477932;

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05-JUL-2004
05-JUL-2004
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
      Submitted (DEC-2001) to the EMBL; BC018747; AAH18747.1; InterPro; IPR003599; Ig. InterPro; IPR007110; Ig-like InterPro; IPR003597; Ig_c1.
                                                                                                                            SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
Strausberg R.;
Submitted (DEC-2001) to
                                                                                                                                                                                                                                                                                  and mouse cDNA sequences Proc. Natl. Acad. Sci. U
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Homo sapiens (Human).
Eukaryota; Metazoa; C
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AAH18747;
02-MAR-2004
02-MAR-2004
SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

TISSUB-Primary B-Cells;

TISSUB-Primary B-Cells;

MEDLINE-22388257; PubMed=12477932;

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Hopkins R.F., Jordan H., Moore T., Max B.J., Wang J., Hong L.,

Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-2004
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27, 27,

Created)

470

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Last annotation update)

update)

Craniata; Ve Catarrhini;

Vertebrata; Euteleostomi;

Hominidae;

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InterPro; IPR003596; Ig_v.

pfam; pF07554; C1-set; 3.

pfam; pF07047; Ig; 4.

SMART; SM00409; IG; 2.

SMART; SM00407; IGc1; 3.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIXE; 4.

PROSITE; PS50835; IG_MHC; UNKNOWN_2.

PROSITE; PS00290; IG_MHC; UNKNOWN_2.

Hypothetical protein.

SEQUENCE 470 AA; 51715 MW; 7849556
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                         EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
WQQGNVFSCSVMHEALHNHYTQKSLSLSFGK
                                                            NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRD
                                                                                                                           NSTYRVVS
                                                                                                                                                                                       GPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQY
                                                                                                                                                                                                     GPSVFLFPPKFKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKFREEQY
                                                                                                                                                                                                                                                 SGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLG
                                                                                                                                                                                                                                                                   SGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLG
                                                                                                                                                                                                                                                                                                               SASTKGPSVFFLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IPR003006; Ig_MHC.
IPR003596; Ig_v.
7654; C1-set; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71.1%; ilarity 91.4%; Conservative 10
                                                                                                                           TTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRD
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Pred. No. 5.2e
10; Mismatches
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Best Local Similarity
Matches 412; Conserv
                                                                                                                                                                                                                                                                                                     Q6PI81
Q6PI81;
O5-UUL-2004 (TrEMBLrel. 2:
O5-UUL-2004 (TrEMBLrel. 2:
O5-UUL-2004 (TrEMBLrel. 2:
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"Generation and initial analysis of more than 15,000 full-length human and mouse CDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
                                                                                                                                                                    Hypothetical protein.
Homo sapiens (Human)
Eukaryota, Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Strausberg R.;
Submitted (DEC-2001) to the
EMBL; BC018747; AAH18747.1;
                      SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical SEQUENCE 47
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                                                                                                                          NCBI_TaxID=9606;
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Pred. No. 5.2e-
10; Mismatches
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A Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Guellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robastein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Robastein M.J., WcKernan K.J., Malek J.A., Gunaratne P.H.,
RA Robast S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Tones S.J. Marra M.A.:
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Jones S.J., Marra M.A.;
"Generation and initial analysis of and mouse cDNA sequences.";
Acad. Sci. U.S.A. 99:16/
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TISSUE=Primary
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CPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNMYVDGVEVHNAK
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                                                                                                 YRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

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A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

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A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

A Whiting M., Madan A., Young A.C., Shewchenko Y., Bouffard G.G.,

A Mhakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Jones S.J., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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MEDLINE=22388257; PubMed=12477932;
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Mammalia; Eutheria;
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Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71.1%;
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 2184; DB 2;
Pred. No. 5.3e-132;
3; Mismatches 23;
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Best Local S
Matches 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     L SUBmitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; BX640947; CAB45972.1; -.

R InterPro; IPR003599; Ig.-1ike.

R InterPro; IPR003597; Ig_c1.

R InterPro; IPR003597; Ig_c1.

R InterPro; IPR003596; Ig_wHC.

R InterPro; IPR003596; Ig_w.

R InterPro; IPR003596; Ig_v.

R Ffam; PF07654; C1-set; 3.

R Pfam; SM0400; IGv; 1.

R PROSITE; PS00290; IG-MHC; UNKNOWN_2.

PROSITE; PS00290; IG-MHC; UNKNOWN_2.

SEQUENCE 475 AA; 52043 MW; B7EAE255A26F4B8E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKPZp686G11190.
Name=DKPZp686G11190;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q6MZQ6
Q6MZQ6;
pcMZQ6;
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THE GERMAN HUMAN CDNA CONSORTIUM;
Lauber J., Bahr A., Mewes H.W., Weil B.,
Han M., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                        AVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPA
                                                                                                                                                                                                          LVSVSAASTKGPSVPPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFP
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PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
                                                                                          AVLOSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPA
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89.0%;
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Best Local Similarity
Matches 406; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae;
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02-MAR-2004 (TrEMBLrel. 27, Last sequence up
02-MAR-2004 (TrEMBLrel. 27, Last annotation
Hypothetical protein DKFZp686G11190.
DKFZP686G11190.
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Lauber J., Bahr A., Mewes H.W.,
Han M., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein. SEQUENCE 475 AA; 52043 MW;
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                                 VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
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                                                                                                                                                                                                                                                                                                                        LVSVSAASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTPP
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ilarity 89.0%;
Conservative 1
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                 VMHEGLHNHYTQKSLSLSPGK
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Pred. No. 7e-131;
8; Mismatches 25;
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Best Local S
Matches 406
Q6N089 PRELIMINARY; PRT; 4/2 AA. Q6N089; Q6N089; Q5-JUL-2004 (TrEMBLrel. 27, Created) 05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation updat Hypothetical protein DKPZp686F15220. Name=DKFZp686F15220;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BAC85232;
BAC85232;
02-MAR-2004
02-MAR-2004
02-MAR-2004
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TISSUE-Dermoid tumor;

TISSUE-Dermoid tumor;

Ota T., Nakagawa S., Senoh A., Mizuguchi H., Inagaki H.,

Hata H., Nakagawa K., Mizuno S., Morinaga M., Kawamura M.

Sugiyama T., Irie R., Otsuki T., Sato H., Nishikawa T., S

Kawakami B., Nagai K., Isogai T., Sugano S.;

"NEDO human cDNA sequencing project.";

"NEDO human cDNA sequencing project.";

Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AK129776; BAC85232.1; -.

SEQUENCE 472 AA; 51212 MW; 01BF215F99809164 CRC64;
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
CDNA FLJ26265 fis, clone DMC05516, highly similar to C region.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                      DKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                             PSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTV
                                                                                                                                                                 EEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLP
                                                                                                                                                                                                                  ELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPR
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                                                                                                                                                                                                                                                                                                                 VLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAP
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                                                                                                                                                EEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLP
DKSRWOOGNVFSCSVMHEALHNHYTOKSLSLSPGK
                                                                          PSRDELTKNOVSLTCLVKGFYPSDIAVEWESNGOPENNYKTTPPVLDSDGSFFLYSKLTV
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Primata; C
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Pred. No. 7
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RESULT 10
CAE45781
ID CAE45
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Matches 408
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Wambutt R., Heubner D., Mewes H.W., Weil B., Amic
Pobo G., Han M., Wiemann S.;
Submitted (Aug-2003) to the EMBL/GenBank/DDBJ dat
EMBL; BX640627; CAE45781.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig cl.
InterPro; IPR003597; Ig cl.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_V.
Pfam; PF07654; Cl-set; 3.
Pfam; PF07654; Cl-set; 3.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 13.
SMART; SM00409; IG; 13.
R PROSITE; PS00239; IG_LIKE; 4.
R PROSITE; PS00239; IG_LIKE; 4.
R PROSITE; PS00239; IG_MHC; UNKNOWN_2.
CAE45781 PRELIMINAR
CAE45781;
02-MAR-2004 (TrEMBLrel
02-MAR-2004 (TrEMBLrel
02-MAR-2004 (TrEMBLrel
Hypothetical protein D
DKFZP686F15220.
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Eukaryota; Metazc
Mammalia; Eutheri
NCBI_TaxID=9606;
[1]
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TISSUE=Human rectu
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aryota; Metazoa;
malia; Eutheria;
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(TrEMBLrel.
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472 AA; 5
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Primates;
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         el. 27, Created)
el. 27, Last sequence up
el. 27, Last annotation
DKFZp686P15220.
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Pred. No. 1.8e
16; Mismatches
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RESULT GEOMETRY OF CONTROL OF CON
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Wambutt R., Heubner D., Mewes
Pobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the E
EMBL; BX640627; CAE45781.1;
Hypothetical protein.
SEQUENCE 472 AA; 51724 MW;
                                                                                                                                                                                                                                                                                                                            Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Cho
Mammalia; Eutheria; Pri
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Eukaryota; Metazoa; (
Mammalia; Eutheria; |
NCBI_TaxID=9606;
[1]
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Q6GMW7;
05-JUL-2004
05-JUL-2004
05-JUL-2004
MEDLINE 22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.
Klausmer R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz
                                                                                                                                                                                                                                                                        NCBI_TaxID=9606; [1]
                                                                                                                                                                                                             SEQUENCE FROM N.A.
TISSUE=Spleen;
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Primata; C
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Primates;
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D., Mewes H.W.,
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Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Young A.C., Shevchenko Y., Bouffard G.G.,
Villalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Villalon M., Touchman J.W., Green E.D., Dickson M.C.,
Villalon M., Touchman J.W., Green E.D., Dickson M.C.,
Villalon M., Touchman J.W., Green E.D., Dickson M.C.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Villalon M., Skalska U., Smailus D.E., Schnerch A., Skalska U., Smailus D.E., Skalska U., 
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Best Local :
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Interpro; IPR003597; Ig_cl.
Interpro; IPR003597; Ig_cl.
Interpro; IPR003596; Ig_wc.
Interpro; IPR003596; Ig_v.
Pfam; PF00647; Ig; 4.
Pfam; PF00047; Ig; 4.
SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 3.
SMART; SM00407; IGc1; 3.
SMART; SM00407; IGc1; 3.
SMART; SM00407; IGc1; 4.
PR0SITE; PS00290; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 475 AA; 51987 MW; 2AIFE
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Submitted (JUN-2004) to the
SMBL; BC073782; AAH73782.1;
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                                                       VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                                            REEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTL
                                                                                                                                                                                                                                                                                     PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
                                                                                                                                                                                                                                                                                                               PELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKP
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                                                                                                                                                                                                     REEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPI
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                           VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
                                                                                                              PPSRDELTKNOVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLT
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475 AA; 51987 MW; 2AlFE55D736860F8 CRC64;
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Pred. No. 2.6e-130;
5; Mismatches 26; 1
                           475
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X Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Carninci P., Prange C.,

A Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., MoEwan P.J., McKerman K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Schmutz J., Myers R.M., Butterfield Y.S.,

A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 408
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AAH06402;
02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (APR-2001) to the EMBL; BC006402; AAH06402.1; Hypothetical protein. SEQUENCE 479 AA; 52281 MP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Strausberg R.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jones S.J., Marra M.A.; "Generation and initial analysis
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Mammalia; Eutheria;
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Homo sapiens
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c. Natl. Acad. Sci. U.S.
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   VYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLY
                                                                                              KTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQ
                                                                                                                                                                                                                     PCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNA
                                                                                                                                                                                                                                                                                                       HTEPAVLQSSGLYSLSSVYTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCP
                                                                                                                                                                                                                                                                                                                                             HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCP
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                                                           KTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQ
                                                                                                                                                                                   PCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNA
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Primates;
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88.7%;
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No. 4.2e-130;
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RESULT
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Best Local S
Matches 401
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TISSUE-Human small intestine;
THE GERMAN HUMAN CDNA CONSORTIUM;
THE GERMAN HUMAN CDNA CONSORTIUM;
Bloecker H., Boecher M., Mewes H.W., Weil B., Ami
Pobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ dat
FROBL; BX640833 (CAB45920.1; -.
InterPro; IPR003109; Ig.like.
InterPro; IPR003109; Ig_like.
InterPro; IPR003109; Ig_MHC.
InterPro; IPR003597; Ig_C1.
InterPro; IPR003597; Ig_C1.
R InterPro; IPR003598; Ig_V.
Pfam; PF00654; C1-set; 3.
Pfam; PF0047; ig; 4.
SWART; SW00409; IG; 2.
SWART; SW00409; IG; 2.
SWART; SW00406; IGV; I.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q6MZV7; PRELIMINARY; PRT; 473 AA.
Q6MZV7;
Q5MZV7;
Q5-JUL-2004 (TrEMBLrel. 27, Created)
Q5-JUL-2004 (TrEMBLrel. 27, Last sequence updated)
Q5-JUL-2004 (TrEMBLrel. 27, Last annotation updated)
Q5-JUL-2004 (TrEMBLrel. 27, Last annotation updated)
Mame-DKFZp686C11235;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Verteb
Mammalia; Eutheria; Primates; Catarrhini; Homi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical SEQUENCE 47
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                                                                                                                                                                                                                                                                               SRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
                                              EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPP
                                                                                                             LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
                                                                                                                                                                         LQSSGLYSLSSVVTVPSSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPB
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                                EQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPI EKTI SKAKGQPREPQVYTLPP
                                                                                             LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
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473 AA; 5
                                                                                                                                                                                                                                                                                                                                                                                                         70.1%; Score 2152.5; DB 2; Length 473; llarity 88.3%; Pred. No. 5.5e-130; Conservative 24; Mismatches 24; Indels 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52121 MW; 9476EAE4C0BFC447 CRC64;
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Catarrhini; Hominidae;
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Best Local Similarity
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BAC85444
BAC85444;
02-MAR-2004
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02-MAR-2004 (TrEMBLrel. 27, Last seq
02-MAR-2004 (TrEMBLrel. 27, Last ann
Hypothetical protein DKFZp686C11235.
DKFZP686C11235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bloecker H., Boecher M., Mewes H.W., Weil B., Fobo G., Han M., Wiemann S.; Submitted (AUG-2003) to the EMBL/GenBank/DDBJ EMBL; BX640853; CA045920.1; -. Hypothetical protein.
SEQUENCE 473 AA; 52121 MW; 9476EAE4COBFC44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAE45920
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TISSUE=Human small intestine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPRE
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                                                                                                                                                                                         SRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
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 (TrEMBLrel.
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                                PRELIMINARY;
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Pred. No. 5.5e-1
24; Mismatches
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Last annotation updat
   Created)
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                                468
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.5e-130;
es 24; Indels
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Best Local Similarity
Matches 400; Conservat
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02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
CDNA FLJ27334 fis, clone TMS09201, highly similar to
C region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "NEDO human cDNA sequencing project.";
Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AK130844; BAC85444.1; -.
SEQUENCE 468 AA; 51224 MW; 9280ACAD6817FC20 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalla; Eutheria; Primata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Thymus;
Kawakami B., Sugiyama A., Takemoto M., Suzuki Y., Hata H.,
Nakagawa K., Mizuno S., Morinaga M., Kawamura M., Sugiyama T.,
Irie R., Otsuki T., Sato H., Nishikawa T., Nagai K., Isogai T.,
Sugano S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
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                                                383
443
                        424
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                                                                                                                                                                                                                                                                                                             GPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYS 183
              VFSCSVMHEALHNHYTQKSLSLSPGK 449
                                               LFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYR 303
                                                                                                                                                                                                          LSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVF 243
                                                                                                                                                                                                                                                                                                                                                                  LVESGGGLVKPEGSLRLSCSASGFTFSDYYMGWTRQAPGKGLEWISYISDSGLSQYHAES
                                                                                                    VVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKN
                                                                                                                    VVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKN 363
                                                                                                                                                      LFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYR 322
VESCSVMHEALHNHYTQKSLSLSPGK
                                                                                                                                                                                                                                                                                                                                                                                                                    69.9%; Score 2148; DB 2; Length 46; ilarity 89.7%; Pred. No. 1.1e-129; Conservative 18; Mismatches 28; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 468;
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